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ALIGNMENTS

REFERENCE AUTHORS TITLE RESULT BI770453 COMMENT SOURCE ORGANISM FEATURES KEYWORDS ACCESSION DEFINITION VERSION Locus JOURNAL source Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 799)

NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Collection (MGC)

Unpublished (1999) BI770453 799 bp mRNA linear LST 43-SEF 40 603055418F1 NIH_MGC_122 Homo sapiens cDNA clone IMAGE:5204692 5', Email: cgapbs-r@mail.nih.gov Tissue Procurement: Life Technologies, found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov Plate: LLAM11513 row: d column: 05 Homo sapiens EST mRNA sequence. Contact: Robert Strausberg, Ph.D. BI770453.1 GI:15762031 CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be ıuman. quality sequence stop: 799.
Location/Qualifiers /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="Ib-MAGE:5204692"
/clone_lib-"NIH_MGC_122"
/lab_host="DH10B" /note="Torgan: pooled lung and spleen; Vector: pCMV-SPORT6; Site_1: Not1; Site_2: EcoRV (destroyed); RNA source anonymous pool of 24 week female lung, 16 week female spleen, and 20-22 week male spleens. Library is oligo-dr primed and directionally cloned (EcoRV site is destroyed upon cloning). Average insert size 1.4 kb, insert size Inc. EST 25-SEP-2001

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Pred. No. 5.7e-135;
0; Mismatches 11;
 bp mRNA linear EST 25-SEP-20 sapiens cDNA clone IMAGE:5205132 5',
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Mammalia; Eutheria; Primates; Catar
1 (bases 1 to 730)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mamm
Unpublished (199)
Contact: Robert Strausberg, Ph.D.
                            GGCACCAGGCAGCTAACAGACACAGCCGCCAGCCCCAAACAGCAGCGGCATGGGCAGCGCC
                                                                                                                              GCCCTGCTGGCCCCTGGTTCCGCTGGCCCCCAGATGCCTGGCTGAGACACGCCAGTGGC
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Tissue Procurement: Life Technologies, Inc.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
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Pred. No. 3.3e-122;
0; Mismatches 11;
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NIH-MGC http://mgc.nci.nih.gov/.
National institutes of Health, M
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                                                                                                                                                                                                                                                                                                                                                                                                       Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP
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                                                          /db_xref="taxon:9606"
/clone="nMAGE:4429413"
/clone="lib="NIH_MGC_91"
/tissue_type="adenocarcinoma, cell line"
/tissue_type="adenocarcinoma, cell line"
/tab_host="DH10B (phage-resistant)"
/note="Organ: prostate; Vector: pCMV-SPORT6; Site_1: NotI;
Site_2: SalI; Cloned unidirectionally; oligo-dT primed.
Average insert size 1.4 kb. Library enriched for
full-length clones and constructed by Life Technologies.
Note: this is a NIH_MGC_Library."
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AI745526
      Mammalia; Eutheria; Primates; Catarrhini; How
1 (bases 1 to 721)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anat
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NCI_CGAP_Pr28 Homo sapiens cDNA clone IMAGE:2317050
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Insert Length: 826 Std Error: 0.00
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Tissue Procurement: Michael J. Brownstein, M.D., Ph.D., Michael
Emmert-Buck, M.D., Ph.D.
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High quality sequence stop: 448.
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/db_xref="taxon:9606"
/clone="IMAGE:2317050"
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Pred. No. 2e-112;
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CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Onpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-rémail.nih.gov
Tissue Procurement: Dr. James Lupski
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National Cancer Institute, Cancer Genome Anatomy
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                                              /note="Vector: PCMV-SPORT6 (Life Technologies); Site_1: NotI; Site_2: SalI; cDNA made by oligo-dT priming. Directionally cloned using the following adaptors: 5'-TCGACCACCGCCGC3' and 5'-GACTAGTTCTAGATCGCGAGCGGCCGCCT[15]-3'. Size selected > 1 kb for average insert length 1.87 kb. This is a primary library, non-amplified. Library constructed by Life Technologies and donated by J. Lupski, M.D./Ph.D. (Baylor College of Medicine) and is available through Life Technologies."
                                         Technologies.
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/db_xref="taxon:9606"
/clone="IMAGE:3394614"
/clone_lib="Lupski_sciatic_nerve"
                                                                                                                                                                                                                                                           /tissue_type="sciatic nerve"
/dev_stage="adult, 70 yr"
/lab_host="DH10B"
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1 (bases 1 to 843)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Ge
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs r@mail.nih.gov
Tissue Procurement: DCTP/DTP
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         CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLN
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information
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           taataaagatactagagaactg
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                                                                  tggtgcagggagacatctgcacccctcagttgggcagccaggagtgcccccggggaatgga
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                                                   GGTGCAAGGGAGACATCTGCACCCCTGAGTTGGGCAGCCAGGAGTGCCCCCGGGAATGGA
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Plate: LLAM10182 row
High quality sequence
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/db_xref="taxon:9666"
/clone="IMAGE:4429712"
/clone=lib="NIH_MGC_91"
/fissue_type="adenocarcinoma, cell line"
/lab_host="DH10B (phage-resistant)"
/note="organ: prostate; Vector: pcMV-SPORT6; Site_1: NotI;
Site_2: SalI; Cloned unidirectionally; oligo-dT primed.
Average insert size 1.4 kb. Library enriched for
full-length clones and constructed by Life Technologies.
Note: this is a NIH_MGC Library."

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Pred. No. 3.4e-102;
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                                                                  cgccctcaggggcctctctcctgcctgccctgccttcagccaggccctgagatgggggaaa 1507
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                                                                                                                              CAGCGCCTCGTCTACCAGTTCGTGCACCCCATCTGAGTGCCTGGCCCAGGGCCTGAAACC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Michael J. Brownstein,
Emmert-Buck, M.D., Ph.D.
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Unpublished (1997)
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BF476482.1
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Seq primer: -40Up from Gibco
High quality sequence stop: 464.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         106
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/db_xref="taxon:9606"
/clone="IMAGE:3256020"
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/sex="male"
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97.0%;
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Pred. No. 8.9e-101;
0; Mismatches 19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Robert Strausberg, Ph.D
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unpublished (1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NIH-MGC http://mgc.nci.nih.gov/
National Institutes of Health, N
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601335410F1 NIH_MGC_39
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                148
         /tissue_type="adenocarcinoma"
/lab_host="DH10B (phage-resistant)"
/lab_host="DH10B (phage-resistant)"
/lab_host="DH10B (phage-resistant)"
/note="Organ: pancreas; Vector: pOTB7; Site_1: XhoI;
Site_2: EcoRI; cDNA made by oligo-dT priming.
Directionally cloned into EcoRI/XhoI sites using the
following 5' adaptor: GGCACGAG(G). Library constructed
by Ling Hong in the laboratory of Gerald M. Rubin
(University of California, Berkeley) using ZAP-cDNA
synthesis kit (Stratagene) and Superscript II RT (Life
Technologies)."

18 a 316 c 319 g 150 t
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                                                                                                                                                                                                      /clone="IMAGE:3689402"
/clone_lib="NIH_MGC_39"
                                                                                                                                                                                                                                         /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                        ocation/Qualifiers
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al Similarity 95.0%;
663; Conservative
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BE871340
BE871340.1 GI:
                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I Mammalla; Eutheria; Primates; Catarrhini; Hominidae; I (bases 1 to 945)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collect
Unpublished (1999)
Contact: Robert Strausberg,
Email: cgapbs-r@mail.nih.go
                                                                Ното
                                                                                                              BE871340
601449457F1 NIH_MGC_65
                                                                sapiens
                                                                                        GI:10320116
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Plate: LLAM9577 row: a column: 15
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/lab_host="DH10B (phage-resistant)"
/note="Organ: colon: Vector: pcMV-SPORT6; Site_1: NotI;
/note="Organ: colon: Vector: pcMV-SPORT6; Site_1: Oligo d'
Site_2: SalI; Cloned unidirectionally. Primer: Oligo d'
Average insert size 1.8 kb. Library constructed by Li
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/db_xref="taxon:9606"
/clone="IMAGE:3853334"
/clone_lib="NIH_MGC_65"
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1 (bases 1 to 625)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Robert Strausberg, Ph.D.
Email: cgapbs-remail.nih.gov
Tissue Procurement: Michael J. Brownstein, M.D.,
Emmert-Buck, M.D., Ph.D.
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Unpublished (1997)
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National Cancer Institute, Cancer Genome Anatomy
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/db_xref="taxon:9606"
/clone="IMAGE:3647751"
/clone_lib="NCI_CGAP_Pr28"
/sex="male"
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                                                                                                                                                                                                                                                                                            /note-"Organ: Prostate; Vector: pT7T3D-Pac (Pharmacia)
/note-"Organ: Prostate; Vector: pT8T3D-Pac (Pharmacia)
with a modified pOlylinker; Plasmid DNA from the
normalized library NCI_CGAP_Pr22 was prepared, and ss
circles were made in vitro. Following HAP purification,
this DNA was used as tracer in a subtractive hybridization
reaction. The driver was PCR-amplified cDNAs from a pool
of 5,000 clones made from the same library (cloneIDs
98508-986759, l101192-1101959, and 1217928-1220615).
Subtraction by Bento Soares and M. Fatima Bonaldo. "
subtraction by Bento Soares and M. Fatima Bonaldo."
a 174 c 202 g 135 t 1 others
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                'lab_host-"DH10B"
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NCI_CGAP_Pr28 Homo sapiens cDNA clone IMAGE:3647751 3'
TR:095238 099238 ETS TRANSCRIPTION FACTOR PDEF. ;, mRNA
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Pred. No. 4.9e-97;
0; Mismatches 17;
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                                                                                                            http://image.lln1.gov
Plate: LLCM1876 row: e column:
High quality sequence stop: 578.
Location/Qualifiers
                                                                                                                                                                                                                                           Contact: Robert Strausberg, |
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                    NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
Unpublished (1999)
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                                                                                                                                                                                     CDNA Library Preparation: Ling Hong/Rubin Laboratory CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLN DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information
/clone="IMAGE:5420913"
/clone=lib="NIH_MGC_110"
/tissue_type="ductal carcinoma, cell line"
/tab_host="DH10B (phage-resistant)"
/note="Organ: pancreas; Vector: pOTB7; Sit
                                                                       /organism="Homo sapiens"
/db_xref="taxon:9606"
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            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; 1 (bases 1 to 899) NIH-MCC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collections (1999)
                                                                                                                                BG328411 899 bp m
602427749F1 NIH_MGC_15 Homo sapiens
mRNA sequence.
BG328411
BG328411 GI:13134849
                                                                                            Homo sapiens
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  Contact:
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Directionally cloned into EcoRI/XhoI sites using the following 5 adaptor: GGCACGAG(G). Library constructed by Ling Hong in the laboratory of Gerald M. Rübin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies).

Note: this is a NIH_MGC Library."
  Robert
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98.3%;
 Strausberg, Ph.D
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Pred. No. 2.1e-96;
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ggagtcccagtccacccgccacgcccgagcagggcctgtccgccttcgacctctcctact
                                                                        gtcgcggacaggcttggagaa-ggcggcagcggggcagtgggtctcgagagacggggact
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                                                                                                                                                                                                           CTGGGCACCAGGCAGCTAACAGACACAGTCG-CAGCCCAAACAGCAGCGGCATGGGCAGC
                                                                                                                                                                                                                             ctgggcaccaggcagctaacagacacagccgccagcccaaacagcagcagcggcatgggcagc
                                                                                                                                                                                                                                                                               CCAGCCCTGCTGGCTCCGGTTCCGCTGGCCCCCCAGATGCCTGGCTGAGACACGCCAGT
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Plate: LLCM1234 row: b column:
High quality sequence stop: 602.
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Tissue Procurement: ATCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            163
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/lab_host="DH10B (phage=resistant)"
/lab_host="Drgan: colon; Vector: poTB7; Site_1: XhoI; Site_2:
EcoRI: cDNA made by oligo-dT priming. Directionally
cloned into EcoRI/XhoI sites using the following 5'
adaptor: GGCACGAG(G). Size-selected >500bp for average
insert size 1.8kb. Library constructed by Ling Hong in
the laboratory of Gerald M. Rubin (University of
Callfornia, Berkeley) using ZAP-cDNA synthesis kit
(Stratagene) and Superscript II RT (Life Technologies)"
63 a 346 c 277 g 113 t
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/clone_lib="NIH_MGC_15"
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/db_xref="taxon:9606"
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Pred. No. 4.5e-92;
0; Mismatches 92;
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RESULT 13
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ns66c07.s1
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CDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www-blo.lln1.gov/bbrp/image/image.html
Insert Length: 734 Std Error: 0.00
Seq primer: -40m13 fwd. ET from Amersham
High quality sequence stop: 393.
Location/Qualifiers
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Eukaryota; Metazoa; Chordata; Catarrhini; Hominidae;
Mammalla; Eutheria; Primates; Catarrhini; Hominidae;
1 (bases 1 to 606)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Pro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              similar to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Michael J. Brownstein,
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
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                      104
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                  /note-*Organ: prostate; Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; 1st strand cDNA was prepared from normal prostate bulk tissue, and was then primed with a Not I - oligo(dT) primer. Double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library is normalized, and was constructed by Bento Soares and M. Fatima Bonaldo. " a 172 c 204 g 126 t
                                                                                                                                                                                                                                                                     /organism-"Homo sapiens"
/db_xref-"taxon:9606"
/clone-"IMAGE:1188588"
/clone_lib-"NCI_CGAP_Pr22"
                                                                                                                                                                                                            /tissue_type="normal prostate"
/lab_host="DH10B"
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SW:ETS4_DROME
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Homo sapiens cDNA clone IMAGE:1188588 3'
P29775 DNA-BINDING PROTEIN D-ETS-4; mRNA
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RESULT 14
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                                                                                                                                                                          sequence.
AW130657
         Tumor Gene Index
(Inpublished (1997)
(Contact: Robert Strausberg, Ph.D.
Email: cgapbs-remail.nih.gov
Tissue Procurement: Christopher Moskaluk,
Emmert-Buck, M.D., Ph.D.
                                                                        Eukaryota; Metazoa; Chordata; Craniata; Verte
Mammalia; Eutheria; Primates; Catarrhini; Hon
1 (bases 1 to 652)
NCI-GGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anau
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cDNA Library Preparation: Life Technologies,
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Pred. No. 4.5e-92;
"" amatches 14;
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TRANSCRIPTION FACTOR PDEF. ;, mRNA
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                                                                            Anatomy
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DNA Sequencing by: Washington University Genome
Clone distribution: NCI-CGAP clone distribution
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
Seq primer: -40UP from Gibco
AA662204
ns66c07.rl NCI_CGAP_Pr22
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/Clone_lib="NCI_CGAP_Ut3"
/tissue_type="poorly-differentiated endometrial
adenocarcinoma, 2 pooled tumors"
/lab_bost="DH10B"
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/db_xref="taxon:9606"
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AA662204
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 National Cancer Institute, Cancer Tumor Gene Index Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       similar to SW:ETS4_DROME P29775 DNA-BINDING PROTEIN D-ETS-4 ;, mRNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Robert Strausberg, Ph.D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
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/sex="male"
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/clone="IMAGE:1188588"
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Sequence 3339, Ap	Sequence 23683, A	Sequence 2401, Ap	~ `	Sequence 824, App	Sequence 955, App	Sequence 1693, Ap	Sequence 1716, Ap	•	Sequence 1610, Ap	ς.				Sequence 627, App			Sequence 327, App	Sequence 369, App	Sequence 920, App

ALIGNMENTS

GENERAL INFORMATION: APPLICANT: ROSETTA Inpharmatics ITITLE OF INVENTION: Diagnosis and Prognosis of FILE REFERENCE: 9301-175-228 CURRENT APPLICATION NUMBER: PCT/US02/18947 CURRENT FILING DATE: 2002-06-14 PRIOR APPLICATION NUMBER: 60/380,770 PRIOR APPLICATION NUMBER: 60/380,770 PRIOR FILING DATE: 2002-05-14 NUMBER OF SEQ ID NOS: 2699 SEQ ID NO 1406 LENGTH: 1894 TYPE: DNA ORGANISM: Homo sapiens PUBLICATION INFORMATION: DATABASE ENTRY DATE: 2001-06-18 PCT-US02-18947-1406 Best Local Similarity 99.3 Matches 1879; Conservative 98.7%; 99.2%; 0 Score 1870; Pred. No. 0; Pred. No. 0; 0; Mismatches DB of. Breast Cancer 15; Length 1894; Indels Patients 0 Gaps

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APPLICANT: Linsley, Peter
APPLICANT: Mao, Mao
APPLICANT: Mao, Mao
APPLICANT: Van 't Veer, Laura
APPLICANT: Van 't Veer, Laura
APPLICANT: Van 't Veer, Marc
APPLICANT: Bernards, Rene
TITLE OF INVENTION: Diagnosis and Prognosis of
FILE REFERENCE: 9301-175-999
CURRENT APPLICATION NUMBER: US/10/172,118
CURRENT FILING DATE: 2002-06-14
PRIOR APPLICATION NUMBER: 60/380,770
PRIOR FILING DATE: 2002-05-14
NUMBER OF SEQ ID NOS: 2699
SEQ ID NO 1406
LENGTH: 1894
TYPE: DNA
ORGANISM: Homo sapiens
PUBLICATION INFORMATION:
DATABASE ACCESSION MUMBER: NM_012391
DATABASE ENTRY DATE: 2001-06-18
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APPLICANT: Dai, Hongyue APPLICANT: He, Yudong
                                                                            Query Match
Best Local Similarity
Matches 1879; Conser
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RESULT

US-10-157-031-83

US-10-157-031-83

Sequence 83, Application US/10157031

GENERAL INFORMATION:

APPLICANT: Baranova, A. V.

APPLICANT: Kozlov, A. P.

APPLICANT: Kozlov, A. P.

APPLICANT: Krukovskaya, L. L.

TITLE OF INVENTION: In slico screening

FILE REFERENCE: 2760-103

CURRENT APPLICATION NUMBER: US/10/157,03

CURRENT ETLING DATE: 2002-05-30

NUMBER OF SEQ ID NOS: 415

SOFTWARE: Patentin version 3.1

SEQ ID NO 83

LENGTH: 1942

TYPE: DNA

ORGANISM: Homo sapiens

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US-10-102-806-215
; Sequence 215, Application
; GENERAL INFORMATION:
; APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Protein:
FILE REFERENCE: PAL03P1C1
CURRENT APPLICATION NUMBER: US/10/102,806
CURRENT FILING DATE: 2002-03-22
PRIOR APPLICATION NUMBER: 09/925,298
PRIOR FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: PCT/US00/05881
PRIOR FILING DATE: 2000-03-08
PRIOR APPLICATION NUMBER: 60/124,270
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; LOCATION: (1075)
; OTHER INFORMATION: n equals
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NUMBER OF SEQ ID NOS: 846
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 215
LENGTH: 1087
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Pred. No. 7.8e
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; NAME/KEY: misc_feature
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APPLICANT: Shyjan, Andrew W.
APPLICANT: Turner, Christopher M.
TITLE OF INVENTION: GENES EXPRESSED IN LUNG C
FILE REFERENCE: PA-0040 US
CURRENT APPLICATION NUMBER: US/09/971,429B
CURRENT FILING DATE: 2001-10-04
PRIOR APPLICATION NUMBER: 60/239,024
PRIOR APPLICATION NUMBER: 60/239,024
PRIOR APPLICATION NUMBER: 60/239,024
PRIOR FILING DATE: 2000-04-10
                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 796; Conserv
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TYPE: DNA
ORGANISM: Homo:
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gcgcctcgtctaccagttcgtgcaccccatctgagtgcctggcccagggcctgaaacccg
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llarity 98.9%;
Conservative
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Pred. No. 1.6e
0; Mismatches
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les 9;
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APPLICANT: Chenchik, Alex
APPLICANT: Bibliashvilli, Robert
TITLE OF INVENTION: uncleic acid arrays
FILE REFERENCE: CLON-006
CURRENT APPLICATION NUMBER: US/09/053,375B
CURRENT FILING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 1543
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 626
SEQ ID NO 626
LENGTH: 2286
TYPE: DNA
ORGANISM: Mus musculus
US-09-053-375B-626
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Best Local Similarity
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                           ccggaagccagacatctcccagcgcctcgtctaccagttcgtgcaccccatctgagtgcc 1428
                                                         caacatgaattacgacaagcttagccgggccttgcggtactactatgataagaatatcat
                                                                          cgccatgaactgcgacaagctgagccgctccatccgccagtcttacaagaagggcatcat 1368
                                                                                                                  gttcaagttggtggatgcagaggaggtggcccggctatggggactgcgcaagaacaagac
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-gtgagcggccagaagtttgtctacaagtttgtgtcctacccagaggttgc
                                                                                                                                                                                                                                                       4.2%;
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Pred. No. 2.6e-06;
0; Mismatches 111;
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GENERAL INFORMATION:

APPLICANT: COTIXA COPPORATION

APPLICANT: Stolk, John A.

APPLICANT: XL, Jiangchun

APPLICANT: XL, Jiangchun

APPLICANT: Chenault, Ruth A.

APPLICANT: Chenault, Ruth A.

APPLICANT: Chenault, Ruth A.

APPLICANT: Compositions and METHODS FOR TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER FILE REFERENCE: 20121.561PC

CURRENT APPLICATION NUMBER: PCT/US01/43704

CURRENT FILING DATE: 2001-11-19

NUMBER OF SEQ ID NOS: 2606

SOFTWARE: COTIXA INVENTION DISCLOSURE DATABASE

SOFTWARE: COTIXA INVENTION DISCLOSURE DATABASE

SOFTWARE: DIAA

ORGANISM: HOMO Sapiens

PCT-1501-44704-2700
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PCT-US01-43704-2290/c
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                                                                                                                                                                                                                                                         Sequence 74, Application PC/TUS0207826
GENERAL INFORMATION:
APPLICANT: Millennium Pharmaceuticals, Inc. et al.
TITLE OF INVENTION: Nucleic Acid Molecules and Proteins For The Identification,
TITLE OF INVENTION: Assessment, Prevention, and Therapy of Ovarian Cancer
FILE REFERENCE: MRI-030PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 148; Conserv
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                                                                PRIOR APPLICATION NUMBER: 60/276,025
PRIOR FILING DATE: 2001-03-14
PRIOR APPLICATION NUMBER: 60/325,149
PRIOR FILING DATE: 2001-09-27
PRIOR APPLICATION NUMBER: 60/276,026
PRIOR FILING DATE: 2001-03-14
                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: PCT/US02/07826
CURRENT FILING DATE: 2002-03-14
                                          PRIOR APPLICATION NUMBER: 60/324,967
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FILING DATE:
APPLICATION
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                   2001/09/26
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Pred. No. 2.8e-05;
0; Mismatches 100;
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US-10-097-340-74
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APPLICATION NUMBER: 60/276,026
PRIOR APPLICATION NUMBER: 60/276,026
PRIOR APPLICATION NUMBER: 60/276,026
PRIOR APPLICATION NUMBER: 60/276,026
PRIOR APPLICATION NUMBER: 60/325,149
PRIOR APPLICATION NUMBER: 60/325,149
PRIOR FILING DATE: 2001-03-14
PRIOR APPLICATION NUMBER: 60/325,149
PRIOR APPLICATION NUMBER: 60/325,149
PRIOR APPLICATION NUMBER: 60/326,026
PRIOR APPLICATION NUMBER: 60/376,026
PRIOR APPLICATION NUMBER: 60/376,026
PRIOR APPLICATION NUMBER: 60/376,026
PRIOR APPLICATION NUMBER: 60/374,026
PRIOR APPLICATION NUMBER: 60/374,026
PRIOR APPLICATION NUMBER: 60/374,026
PRIOR APPLICATION NUMBER: 60/374,967
PRIOR PILING DATE: 2001/09/26
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APPLICANT:
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Best Local :
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APPLICANT:
APPLICANT:
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PRIOR APPLICATION NUMBER: 60/325,102
PRIOR FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: 60/323,580
PRIOR FILING DATE: 2001-09-19
NUMBER OF SEQ ID NOS: 363
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ORGANISM: Homo sapiens
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Local Similarity 59.0%;
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Karen GLATT
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Rachel E. MEYERS
Michael MORRISEY
                                                                                                                                                                                                                                                                                                                                                                                Gordon B. MILLS
Robert C. BAST, Jr.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sebastian HOERSCH
Shubhangi KAMATKAR
                                                                                                                                                                                                                                                                                                                                                                                                                                           Ami SEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Manjula GANNAVARAPU
                                                                                                                                                                                                                                                                                                                                         Rosemarie SCHMANDT
                                                                                                                                                                                                                                                                                                                                                               Karen LU
                                                                                                                                                                                                                                                                                                                                                                                                                        Peter VEIBY
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Peter OLANDT
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Pred. No. 2.8e-05;
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TITLE OF INVENTION: NOVEL CONTIGS OBTAINED FROM VARIOUS CDNA
TITLE OF INVENTION: LIBRARIES
FILE REFERENCE: 20411-752CON1
CURRENT APPLICATION NUMBER: US/09/919,002
CURRENT APPLICATION NUMBER: APPLICATION NUMBER: 09/359,922
PRIOR APPLICATION NUMBER: APPLICATION NUMBER: 09/359,922
PRIOR FILING DATE: FILING DATE: 1999-07-22
PRIOR APPLICATION NUMBER: APPLICATION NUMBER: US 09/034,341
PRIOR FILING DATE: FILING DATE: 1998-02-13
NUMBER OF SEQ ID NOS: 13203
SOFTWARE: FASTSEQ for Windows Version 3.0
SEO ID NO 13047
LENGTH: 2457
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PRIOR APPLICATION NUMBER: 60/325,102
PRIOR APPLICATION NUMBER: 60/323,580
PRIOR APPLICATION NUMBER: 60/323,580
PRIOR FILING DATE: 2001-09-19
NUMBER OF SEQ ID NOS: 363
SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 74
LENGTH: 1907
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                                                                                                                                                                                                                                                   ; ORGANISM: Homo sapiens US-09-919-002-13047
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                                                                                                                                                              Query Match
Best Local Similarity
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Best Local Similarity 59.0%;
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1219 cattaggtggctcaacaaggagaagggcatcttcaaaaattgaggactcagcccaggtggc 1278
                                                                                1159 catccacctgtggcagttcctcaaggagttgctactcaagccccacagctatggccgctt 1218
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                                                           940 cacccacctgtgggagttcatccgggacatcctcatccacccggagctcaacgagggcct 999
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Pred. No. 2.8e-05;
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Pred. No. 2.8e-05;
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CCURRENT APPLICATION NUMBER: US/10/099,926
CURRENT FILING DATE: 2002-03-17
NUMBER OF SEQ ID NOS: 1982
SOFTWARE: FBASTSQ for Windows Version 4.0
SEQ ID NO 1680
LENGTH: 2642
TYPE: DNA
ORGANISM: Homo sapiens
US-10-099-926-1680
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APPLICANT: King, Gordon E.
APPLICANT: Meagher, Madeleine Joy
APPLICANT: Xu, Jiangchun
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Best Local
                                                                                                                                                         APPLICANT: Lille, James
TITLE OF INVENTION: COMPOSITIONS, KITS, AND METHODS FOR
TITLE OF INVENTION: IDENTIFICATION, ASSESSMENT, PREVENTION, AND THERAPY OF
TITLE OF INVENTION: OVARIAN CANCER
FILE REFERENCE: MRI-006A
CURRENT EPPLICATION NUMBER: US/09/654,788A
CURRENT FILING DATE: 2000-09-01
                                                                                  CURRENT FILING DATE: 2000-09-01
PRIOR APPLICATION NUMBER: 60/152,547
PRIOR FILING DATE: 1999-09
PRIOR APPLICATION NUMBER: 60/190,347
PRIOR FILING DATE: 2000-03-16
PRIOR FILING DATE: 2000-03-16
                                                                                                                                                                                                                                                                   APPLICANT: Lee, John APPLICANT: Thompson, APPLICANT: Lillie,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY TITLE OF INVENTION: AND DIAGNOSIS OF COLON CANCER FILE REFERENCE: 210121.547C2
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                                                APPLICATION NUMBER: 60/191,321 FILING DATE: 2000-03-21
                APPLICATION NUMBER: 60/208,382 FILING DATE: 2000-05-31
APPLICATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                tgagccgctccatccgccagtcttacaagaagggcatcatccggaag 1375
                                                                                                                                                                                                                                                                                                                                                                                                                                tgagccgctcgctccgatactattatgagaaaggcatcatgcagaag 1351
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87; Conserv
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Jiang, Yuqiu
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NUMBER:
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81.3%;
60/220,467
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Pred. No. 2.8e-05;
0; Mismatches 20;
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GENERAL INFORMATION:
APPLICANT: FUJINAGA, Kei
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Best Local
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                                                                                  APPLICATION NUMBER: PCT/JP96/00016 FILING DATE: 09-JAN-1996 ATTORNEY/AGENT INFORMATION:
             REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                              CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/167,485
FILING DATE: 13-Jun-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HIGASHINO, Fumihiro TITLE OF INVENTION: CANCER CONTROL NUMBER OF SEQUENCES: 5
                                                                                                                                                                                                               PRIOR APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ctacaagtttg 1229
                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0,
                                              NAME: BROWDY, ROGER L. REGISTRATION NUMBER: 2
                                                                                                                                      FILING DATE: <Unknown>
APPLICATION NUMBER: 08,
FILING DATE: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                              STATE: D.C
                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: Washington
                                                                                                                                                                                            APPLICATION NUMBER: US/09/602,868
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                                                                                                                                                                                                                                                                                                                                                                                            ZIP:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: 624 Ninth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: BROWDY AND NEIMARK, P.L.L.C. STREET: 624 Ninth Street N.W., Ste. 300
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1159 catccacctgtggcagttcctcaaggagttgctactcaagccccacagctatggccgctt 1218
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                                   catccgccagtcttacaagaaggcatcatccggaagccagacatctcccagcgcctcgt 1398
catgag---gtactactacaaacgggagatcctggaacgggtggatggccggcgactcgt
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Pred. No. 2.8e
0; Mismatches
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     1218
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TELEFAX: (202) 737-3528
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:

LENGTH: 2064 base pairs

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CURRENT APPLICATION NUMBER: US/10/099,926; CURRENT FILING DATE: 2002-03-17; NUMBER OF SEQ ID NOS: 1982; SOFTWARE: FASTSEQ for Windows Version 4.0; SEQ ID NO 1788; LENGTH: 2333; TYPE: DNA ORGANISM: Homo sapiens
US-10-099-926-1788
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APPLICANT: King, Gordon E.
APPLICANT: Meagher, Madeleine Joy
APPLICANT: Xu, Jiangchun
APPLICANT: Secrist, Heather
APPLICANT: Jiang, Yugiu
                                                                                                                                                                                                                    Query Match 3.9%;
Best Local Similarity 74.6%;
Matches 94; Conservative
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Best Local Similarity 74.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY TITLE OF INVENTION: AND DIAGNOSIS OF COLON CANCER FILE REFERENCE: 210121.547C2
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1428
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                                                                                          gccatgaactgcgacaagctgagccgctccatccgccagtcttacaagaagggcatcatc 1369
cagaag
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                                                                    gccatgaattacgacaagctgagccgctcgctccgatactattatgagaaaggcatcatg 1427
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STRANDEDNESS: single
TOPOLOGY: linear
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LOCATION: 1...
1433
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Pred. No. 3e-05;
""ematches 32;
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Pred. No. 3e-05;
0; Mismatches 32;
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Search completed: August Job time: 9463 sec

6, 2002,

RESULT

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CURRENT APPLICATION NUMBER: PCT/US02/12378
CURRENT FILING DATE: 2002-04-15
NUMBER OF SEQ ID NOS: 627
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 282
LENGTH: 502
TYPE: DNA
ORGANISM: Homo sapiens
PCT-US02-12378-282
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Best Local S
Matches 125
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APPLICANT:
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF BREAST CANCER
FILE REFERENCE: 210121.47003PC
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APPLICANT:
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Local Similarity 59.5%;
hes 125; Conservative
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                        catccgccagtcttacaagaagggcatcat 1368
                                                                                                     ccggctgtggggcatccgcaagaaccgtcccgccatgaactgcgaccagctgagccgctc 1338
catgaggtactactacaaacgggagatcct 478
                                                                              ccaactatggggccaaaagaaaagaacagcaacatgacctacgagaagctgagccgggc 448
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Jiang, Yuqiu
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Perfect score:
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US-09-439-313-398
US-09-213-767-1
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US-08-16-049-3
US-08-746-789A-1
US-08-343-443B-3
US-08-780-835B-1
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US-09-020-956-44
US-09-030-607-44
US-09-030-6313-3
US-08-368-281-3
US-08-368-281-3
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; CLONE: US-09-055-113-2
                                                                                                                                                                                                                         APPLICATION NUMBER: US/09/055,113
FILING DATE: Filed Herewith
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILLING DATE:
APPLICATION NUMBER:
ATTORNEY/AGENT INFORMATION:
NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-0501 US
TELEPHONE: 650-855-0555
TELEPHONE: 650-855-0555
                                                                                                                                TELEX:
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 1905 base pair:
                                                                                                                                                                                                                                                                                                                                                                                                                                             OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
                                                      TOPOLOGY: line
IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Bandman, Olga
APPLICANT: Corley, Neil C.
APPLICANT: Guegler, Karl J.
APPLICANT: Lal, Preeti
MMEDILL
LIBRARY: FNC.
1813005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM CONTROL OPERATING SYSTEM:
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CITY: Palo Alto
STATE: CA
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                                                                                            STRANDEDNESS:
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                                     999caccctggacttggtgcccggcggctgaccttggaggagcactcgctggagcaggt
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RESULT 2
US-09-439-313-398
; Sequence 398, Application US; Patent No. 6329505
; GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
; APPLICANT: Dillon, Davin (
APPLICANT: Mitcham, Jennil
; APPLICANT: Harlocker, Susciapper, Susc
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Xu, Jiangchun C.
Dillon, Davin C.
Mitcham, Jennifer I
Harlocker, Susan Lc
Jiang Yugui
Reed, Steven G.
Kalos, Michael
Fanger, Gary
Retter, Mark
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; LCCATION: (1)...(278)
; OTHER INFORMATION: n = A,T,C
US-09-439-313-398
                                                                                                                                                    ; NAME/KEY: CDS
; LOCATION: (316)..(1602)
US-09-213-767-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 1, Application US/09213767 Patent No. 5948680 GENERAL INFORMATION:
                                                    Query Match 4.4%;
Best Local Similarity 59.8%;
Matches 177; Conservative
                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 47
SEQ ID NO 1
LENGTH: 2266
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Best Local Similarity
Matches 263; Conserv
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CURRENT FILING DATE: 1999-11-12
NUMBER OF SEQ ID NOS: 575
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 398
LENGTH: 278
TYPE: DNA
                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/213,767
CURRENT FILING DATE: 1998-12-17
                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Brenda F. Baker
APPLICANT: Lex M. COWSETT
TITLE OF INVENTION: ANTISENSE MODULATION OF ELK-1 EXPRESSION
FILE REFERENCE: RTS-0024
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                                                                                                                                                                                                                           TYPE: DNA
ORGANISM: Homo sapiens
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1166 ctgtggcagttcctcaaggagttgctactcaagccccacagctatggccgcttcattagg 1225
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Day, Craig
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                                                      Score 83.2; DB 2;
Pred. No. 2.5e-09;
0; Mismatches 113;
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                                                                                        TELEFAX: (415) 576-0200
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 2667 base pairs
TYPE: nucleic acid
STRANEDNESS: single
TOPOLOGY: linear
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Patent No. 58561
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/469,412A
FILING DATE: 05-JUN-1995
CLASSIFICATION: 435
CLASSIFICATION: 435
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                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: 01 TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION: NAME: Garrett-Wackowski,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Athanasiou, Meropi A.
APPLICANT: Sgouras, Dionyssios N.
TITLE OF INVENTION: The ERF Genetic Locus and Its Products
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LOCATION:
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CITY: San Francisco
OTHER INFORMATION: /note= "human OTHER INFORMATION: Factor) cDNA"
                                                                                                                                                                                                                                                                                                NAME: Garrett-Wackowski, REGISTRATION NUMBER: 37,3
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                 ERF (ETS2 Repressor
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Query Match Best Local Similarity Matches 150; Conserv

Conservative

4.0%; 59.1%;

Score 75.2; DB 2; Pred. No. 1.3e-07; 0; Mismatches 98;

Length 2667; Indels

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RESULT 5
US-09-021-715-1
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us-09-021-715-1
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Patent No. 6194547
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION UMBER: US/09/021,715
FILING DATE: 10-reb-1998
CLASSIFICATION: <Unknown>
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                            NAME: Garrett-Wackowski, Eugenia
REGISTRATION NUMBER: 37,330
REFERENCE/DOCKET NUMBER: 015280-229000
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
Townsend and Townsend
ADDRESSEE: Townsend and Townsend
ADDRESSEE: Townsend and Townsend
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sgouras, Dionyssios N. TITLE OF INVENTION: The ERF Genetic
                     SEQUENCE DESCRIPTION: SEQ ID NO: 1:
                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
                                                                                                                                              MOLECULE TYPE: CDNA
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                                                                                 NAME/KEY: CDS
LOCATION: 123..1769
                                                                                                                                                                                     LENGTH: 2667 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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                                                           OTHER INFORMATION:
                                                                                                                                                                    TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 94111-3834
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Eighth Floor
                                                             ERF (ETS2 Repressor
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Best Local Similarity 59.1
Matches 150; Conservative
                                                                                            TELEFAX: (202) 737-352
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                     FILING DATE: 07-AUG-1997
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 07-020173
FILING DATE: 08-FEB-1995
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                             FILING DATE: 09-JAN-1996 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Release #1.0, V
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/875,944B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: HIGASHINO, Fumihiro TITLE OF INVENTION: CANCER CONTROL NUMBER OF SEQUENCES: 5
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MOLECULE TYPE:
                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
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                  TYPE: nucleic acid
STRANDEDNESS: sing
TOPOLOGY: linear
                                                                                                                                                                                                    NAME: BROWDY, ROGER L. REGISTRATION NUMBER: 25, REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Washington
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                                                                                                                                                          TELEPHONE:
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                                                                             LENGTH:
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                                                                             2064 base pairs
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                                                                                                                                                            (202) 628-5197
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                                     single
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Pred. No. 1.3e-07;
0; Mismatches 98;
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LENGTH: 2064
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-116-049-3
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US-08-746-789A-1
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GENERAL INFORMATION:
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Best Local Similarity
Matches 94; Conserv
                                                                                                                                                                                              Sequence 1, Application US/08746789A Patent No. 5789200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                      APPLICANT: Ismail Kola, Martin J. Tymms, Christine DeBouck APPLICANT: Ismail Kola, Martin J. Tymms, Christine DeBouck TTTLE OF INVENTION: A No. 5789200el Human ETS Family Member,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/116,049A
CURRENT FILING DATE: 1998-07-15
NUMBER OF SEQ ID NOS: 11
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Hung, Men-Chie TITLE OF INVENTION: HUMAN PEA3 IS A TUMOR SUPPRESSOR FOR CANCER FILE REFERENCE: UTSC:582
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                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKline Beecham Corporation
STRRET: 709 Swedeland Road, P.O. Box 1539
CITY: King of Prussia
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                    COUNTRY:
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19406-0939
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Conservative (
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1..1386
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Pred. No. 1.5e-07;
0; Mismatches 32;
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Pred. No. 1.5e-07;
0; Mismatches 32;
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US-08-343-443B-3
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Best Local Similarity 58.6%;
Matches 147; Conservative
                                                                                                                                                                                                                                                           Sequence 3, Application US/083434443B Patent No. 5968734
                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
              APPLICANT: Thomas, G
APPLICANT: Zucman, J
TITLE OF INVENTION:
                                                                                                                                    APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                    1167
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NAME: William T. Han
REGISTRATION NUMBER: 34,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: NO. 5789
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                        1399 ctaccagttcg 1409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1219 cattaggtggctcaacaaggagaagggcatcttcaaaaattgaggactcagcccaggtggc 1278
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                                                                                                                                                                                                                                                                                                                                                                                                                                     1110 CATGAG---GTACTACAAACGGGAGATCCTGGAACGGGTGGATGGCCGGCGACTCGT 1166
                                                                                                                                                                                                                                                                                                                                                                                                                                                             1339 catccgccagtcttacaagaagggcatcatccggaaggccagacatctcccagcggcctcgt 1398
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1050 CCAACTATGGGGGCAAAAGAAAAAGAACAGCAACATGACCTACGAGAAGCTGAGCCGGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              930 CACCCACCTGTGGGAGTTCATCCGGGACATCCTCATCCACCCGGAGCTCAACGAGGGCCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: Nucleic Acid
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PATE: No. 5789200ember 15, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                  CTACAAGTTTG 1177
   OF
                                                                                                                 Peter, Martine
Ploougastel, Beatrice
Thomas, Gilles
                                                                                                                                                                      Melot,
                                                                                                                                                                                        Delattre, Olivier
Desmaze, Chantal
                                                                                                     Zucman, Jessica
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NUCLEIC ACID CORRESPONDING TO A GENE OF CHROMOSOME 22 INVOLVED IN RECURRENT CHROMOSOMAL TRANSLATIONS ASSOCIATED WITH THE DEVELOPMENT OF CANCEROUS TUMORS, AND NUCLEIC ACIDS OF FUSION RESULTING FROM SAID TRANSLOCATIONS
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CORRESPONDENCE ADDRESS:

ADDRESSEE:

3: Weiser & Associat 230 South Fifteenth

Associates ifteenth Street

Philadelphia PΑ

19102

USA

P.O. Box 4433

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US-08-780-835B-1
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            Sequence 1, Application US/08780835B
Patent No. 592268B
GENERAL INFORMATION:
APPLICANT: Hung, Mien-Chie
APPLICANT: Xing, Xiangming
TITLE OF INVENTION: PEA3 is a Tumor Suppressor
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
Matches 132; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: PCT/FR
FILING DATE: 19-MAX-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: FR 92/
FILING DATE: 20-MAX-1992
ATTORNEX/AGENT INFORMATION:
NAME: Weiser, Gerard J.
                                                                                                                                                                                                                                                                                                                                                                                                                                       1216 cttcattaggtggctcaaccaaggagaagggcatcttcaaaaattgaggactcagcccaggt 1275
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COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: AEDIT 1.0 DOS text edi
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                           1153 GGCCCTCCGTTATTACTATGATAAAAACATTAT 1185
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TELEFAX: 215-875-8394
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PRIOR APPLICATION DATA:
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TYPE: n
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REGISTRATION NUMBER: 19,763
REFERENCE/DOCKET NUMBER: 989.6121P
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LOCATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: double
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    ADDRESSEE:
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                                                                                                                                                                                                                                                                                            ctccatccgccagtcttacaagaagggcatcat 1368
                                                                                                                                                                                                                                                                                                                                                             99cccggctgtggggcatccgcaagaaccgtcccgccatgaactgcgacaagctgaagccg 1335
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ARNOLD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3.7%; Score 71; DB 2; Le 62.0%; Pred. No. 1.1e-06; n. Mismatches 75;
WHITE AND DURKEE
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RESULT 11
US-09-303-268-1
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                                                                                                                                                                                                                          Sequence 1, Application US/09303268 Patent No. 6172212 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 3.7%;
Best Local Similarity 59.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (512) 474-757
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: USA
COUNTRY: TOSA
TTD: 77210-4433
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REGISTRATION NUMBER: 37,259
REFERENCE, DOCKET NUMBER: UTI
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                    1542 TATTATGAGAAAGGCATCATGCAGAAG 1568
                                                                                                                                                                                                                                                                                                                                                                                                                                                       1289
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CURRENT APPLICATION DATA:
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 10 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH:
                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: ARNOLD,
                                                                                                                                                        Xing, Xiangming TITLE OF INVENTION: PEA3 is a NUMBER OF SEQUENCES: 9
   COMPUTER READABLE MEDIUM TYPE:
                                                                                                                                                                                                          APPLICANT: Hung, Mien-Chie
                                                                                                                                                                                                                                                                                                                                                                      tottacaagaagggcatcatccggaag 1375
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                                                                                                                                                                                                                                                                                                                                                                                                                                        ggcatccgcaagaaccgtcccgccatgaactgcgacaagctgagccgctccatccgccag 1348
                                  COUNTRY: USA
ZIP: 77210-4433
                                                                     CITY: Houston
STATE: Texas
                                                                                                         STREET: P.O. Box 4433
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E: Floppy disk
IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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512) 474-7577
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FORM:
Floppy disk
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                                                                                                                        WHITE AND DURKEE
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Pred. No. 1.6e-06;
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                                                                                                                                                                        Tumor Suppressor
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                                                                          GENERAL INFORMATION:
APPLICANT: Hung, Men-Chie
APPLICANT: Hung, Men-Chie
TITLE OF INVENTION: HUMAN PEA3 IS A TUMOR SUPPRESSOR FOR CANCER CELLS
FILE REFERENCE: UTSC:582
CURRENT FILING DATE: 1998-07-15
CURRENT FILING DATE: 1998-07-15
NUMBER OF SEQ ID NOS: 11
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 1
LENGTH: 2410
TYPE: DNA
ORGANISM: Mus musculus
S-09-116-049-1
Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                Sequence 1, Application US/09116049A Patent No. 6248351
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches 158; Conservative
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Best Local Similarity
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TELEPHONE: (512) 418-3000
TELEPAX: (512) 474-7577
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 2410 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1308 GAAGGGATTGGAGCTTTCCGGGAGGGGCCACCCTACCAGCGCCGGGGTGCCTTACAACTG 1367
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FILING DATE: 10-JAN-1997
ATTORNEY/AGENT INFORMATION:
NAME: Wilson, Mark B.
REGISTRATION NUMBER: 37,259
REGISTRATION NUMBER: UTSC500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 1:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TGGCAGTTTCT---GGTGGCCCTGCTGGATGACCCAACAAATGCTCATTTCATTGCTTGG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0,
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59.2%;
3.7%;
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Score 70.2; DB 4;
Pred. No. 1.6e-06;
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US-09-360-779-1
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APPLICANT: Deneris, Evan S.
APPLICANT: Prodoro, Dmitry V.
APPLICANT: Evodoro, Dmitry V.
APPLICANT: Hendricks, Timothy J.
APPLICANT: CASE-03828
TILE REFERENCE: CASE-03828
FILE REFERENCE: CASE-03828
CURRENT APPLICATION NUMBER: US/09/360,779
CURRENT FILING DATE: 1999-07-26
EARLIER APPLICATION NUMBER: 60/094,264
EARLIER APPLICATION NUMBER: 60/094,264
EARLIER FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 3.4%;
Best Local Similarity 53.1%;
Matches 186; Conservative
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cgtctaccagttcgtgcaccccatctgagtgcctggcccagggcctgaaacccgccctca 1455
                                                                                      agcactgcgctactactacgacaaaaacatcatgagcaaggtgcacggcaagcgctacgc 788
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pred. No. 2.1e-05;
0; Mismatches 157;
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; NAME/KEY: CDS
; LOCATION: (291)..(1700)
US-09-344-579-1
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Patent No. 5856125
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APPLICANT: Lex M. Cowsert
TITLE OF INVENTION: ANTISENSE MODULATION OF ETS-2 EXPRESSION
FILE REFERENCE: RTS-0063
CURRENT APPLICATION NUMBER: US/09/344,579
CURRENT FILING DATE: 1999-06-25
                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Mavrothalassitis
APPLICANT: Blair, Donald G.
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ORGANISM: Homo sapiens
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatii
OPERATING SYSTEM: PC-DOS
                                                                                                                                                                                                                 APPLICANT: Athanasiou, Meropi A.
APPLICANT: Sgouras, Dionyssios N.
TITLE OF INVENTION: The ERF Genetic Locus and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1157 cccatccacctgtggcagttcctcaaggagttgctactcaagccccacagctatggccgc 1216
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CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                          APPLICANT: Fisher, Robert APPLICANT: Beal Jr., Grego
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1548 ggcttacgctactattacgacaagaacatcatccacaagac 1588
                                                                                                         CITY: San Francisco
STATE: California
                                                                      COUNTRY: USA
ZIP: 94111-3834
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
mes 130; Conserv
                                                                                                                                           ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor
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Athanasiou, Meropi A.
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PC-DOS/MS-DOS
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Pred. No.
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; NAME/KEY: 1..2544; LOCATION: 1..2544; OTHER INFORMATION: OTHER INFORMATION: OTHER INFORMATION: US-08-469-412A-6
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Search completed: August Job time: 5741 sec
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INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 2544 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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Best Local Similarity
Matches 123; Conserva
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
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FILING DATE: 05-JUN-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Garrett-Wackowski, Eugenia
REGISTRATION NUMBER: 37,330
                                                                                                         1315 gaactgcgacaagctgagccgctccatccg 1344
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                  6, 2002, 19:52:32
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ERF gene (promoter region and first exon
not included)"
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Maximum DB seq length: 2000000000
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Listing first 45 summaries
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AC008253
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AC008217
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AF106964
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AX048159 Sequence
BC021299 Homo sapi
AB031549 Homo sapi
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AB031549 Homo sapi
BC012648 Mus muscu
AL157372 Human DNA
AX048155 Sequence
AX106617 Sequence
AX10768 Sequence
AX267424 Sequence
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AX267424 Drosophila
AC015374 Drosophila
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AC008213 Drosophila
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AB016193 Homo sapi
AF075706 Gallus ga
AF429315 Homo sapi
AF000672 Homo sapi
AJ249590 Danio rer
Z36939 M.musculus
AX305989 Sequence
X87257 M.musculus
AJ339670 Homo sapi
Y08979 H.sapiens F
Y08976 H.sapiens m
AC097468 Homo sapi
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U84615 Danio rerio
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ALIGNMENTS

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AUTHORS
TITLE
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ORGANISM
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LOCUS
DEFINITION
ACCESSION
VERSION
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                                                                             source
                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Mammalia; Eutheria; Primates; Catarrhir 1 (bases 1 to 1894)
Kaufmann, J. and Xin, H.
Expression of ets-domain proteins in catarent: WO 0070092-A 1 23-NOV-2000; Chiron Corporation (US)
Location/Qualifiers
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Sequence 1 from Patent
AX048148
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416. .1423
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                /note="Human JKETS."
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Patent W00070092.
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WTEHQYRLPPMGKAFQELAGKELCAMSEEQFRQRSPLGGDVLHAHLDIWKSAAWMKER
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900	841 catcaccgcagatcccatggactggagccccagcaatgtgcagaagtggctcctgtggac :	Оy

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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
1 (bases 1 to 3317)
Kaufmann,J. and Xin,H.
Expression of ets-domain proteins in cancer
Patent: WO 0070022-A 12 23-NOY-2000;
Chiron Corporation (US)
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PSPPARTDEQGLSAFYLSYFDMLYPEDSSWAAKAGASSREEPPESPEQCPVIDSQAPA
GSLDLYPGGLTLEEHSLEVQUSMYCGSVLADIETACKLLNITADPMWSPSPSWYQKHL
WTEHQYRLPPMGKAFQELAGKELCAMSEEQFRQRSPLGGDVLHAHLDIWKSAAWMKER
TSPGALHYCASTSEESWTDSEVDSSCSGQPIHLWQFLKELLLRHSYGRFIRWLMKEK
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                                                                                                                                                                                                                                                                                                                                                                    Web site: http://www.nisc.nih.gov/
Contact: nisc_mgc@nhgri.nih.gov/
Shevchenko,Y., Wetherby,K.D., Beckstrom-Sternberg,S.M.,
Benjamin,B., Blakesley,R.W., Bouffard,G.G., Brinkley,C., Brooks,S.,
Benjamin,B., Blakesley,R.W., Bouffard,G.G., Brinkley,C., Brooks,S.,
Dietrich,N.L., Guan,X., Gupta,J., Ho,S.-L., Karlins,E., Legaspi,R.,
Lim,M., Maduro,Q.L., Mastello,C., Mastrian,S.D., McCloskey,J.C.,
McDowell,J., Pearson,R., Snyder,B., Stantripop,S., Thomas,P.J.,
Tiongson,E.E., Touchman,J.W., Tsurgeon,C., Vogt,J.L., Walker,M.A.,
                                                                                                                                                                                                     Clone distribution: MGC clone distribution through the I.M.A.G.E. Consortium/IIML at: Series: IRAL Plate: 39 Row: d Column: 22 This clone was selected for full length sec passed the following selection criteria: ma Location/Qualifiers
1. 1942
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cDNA Library Preparation: Rubin Laboratory
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: National Institutes of Health Intramural
Sequencing Center (NISC),
Gaithersburg, Maryland;
Web site: http://www.nisc.nih.gov/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Submitted (14-JAN-2002) National Institutes of Health, Mammalian Gene Collection (MGC), Cancer Genomics Office, National Cancer Institute, 31 Center Drive, Room 11A03, Bethesda, MD 20892-2590,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NIH-MGC Project URL: http://mgc.nci.nih.gov
Contact: MGC help desk
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/clone="MGC:29592 IMAGE:4546860"
/tissue_type="Colon, adenocarcinoma"
/clone_lib="NIH_MGC_15"
/lab_host="DH10B-R"
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1864	cctggcatggtgcagggagacatctgcacccctcagttgggcagccagggagtgccccc	1805	Qy
1809	CTCTCTGCTCCATCCCCCTGCCTCCCATTC	1750	DЪ
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1744		1685	Qy
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1629	CCAACTGCCCCAGGGGATATGGGTCCTCTGGGGCCTTCGGGACCCTGGGGCAGGGGTGC	1570	рь
1624	caactgccccagggggatatgggtcctctggggccttcgggaccatggggcaggggtg	1565	Qy
1569	AAAACGGGCAGTCTGCTCTGCTCTCTGACCTTCCAGAGCCCAAGGTCAGGGAGGG	1510	Db
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1329	ACGAC	1270	DЪ
1324	caycocaygtggcccggctgtggggcatccgcaagaaccgtcccgccatgaactgcgac	1265	ν.
1269	AGCTATGGCCGCTTCATTAGGTGGCTCAACAAGGAGAAGGGCATCTTCAAAATTGAGGAC	1210	망
1264	gctatggccgcttcattaggtggctcaacaaggagaagggcatcttcaaaattgaggac	1205	Qy
1209	TGCTCCGGGCAGCCCATCCACCTGTGGCAGTTCCTCAAGGAGTTGCTACTCAAGCCCCAC	1150	ДĎ
1204	gctcogggcagcccatccacctgtggcagttcctcaaggagttgctactcaagccccac	1145	Qy
1149	-	1090	Db
1144	ttcactactgtgcctcgaccagtgaggagagctggaccgacagcgaggtggactcat	1085	Qy
1089		1030	Db
1084	cccacctggacatctggaagtcagcggcctggatgaaagagcggacttcacctgggg	1025	Оу
1029	TGCGCCATGTCGGAGGAGCAGTTCCGCCAGCGCTCGCCCCTGGGTGGG	970	Db
1024		965	Qy
969	CACCAATACCGGCTGCCCCCATGGCCAAGGCCTTCCAGGAGCTGGCCGGGCAAGGAGCTG	910	Db
964	accaataccggctgccccccatgggcaaggccttccaggagctggcgggcaaggagctg	905	Qy
909		850	DЪ
904	ccgcagatcccatggactggagccccagcaatgtgcagaagtggctcctgtggacagag	845	Qy

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Submitted (25-AUG-1999) Masami Nozaki, Osaka University, Institute for Microbial Diseases; 3-1, Yamadaoka, Suita, Osaka 565-0871, Japan (F-mail.mnozaki@biken.osaka-u.ac.jp, Tel:81-6-879-8338,
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Mammalia; Eutheria;
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                                                                                                                                                                                                                                     /gene="PSE"
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GIFKIEDSAQVARLWGIRKNRPAMNYDKLSRSIRQYYKKGIIRKPDISQRLVYQFVHP
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                    TATIGGCCGCTTCATTAGGTGGCTCAACAAGGAGAAGGGCATCTTCAAAATTGAGGACTCA
                                                            TCCGGGCAGCCCATCCACCTGTGGCAGTTCCTCAAGGAGTTGCTACTCAAGCCCCCACAGC
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Contact: MGC help desk
Email: cgapbs-remail.nih.gov
Tissue Procurement: Jeffrey E. Green, M.D.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium
DNA Sequencing by: Institute for Systems Biology
                                                                                                                               Direct Submission
Direct Submission
Submitted (15-AUG-2001) National Institutes of Health, Mammalian Submitted (15-AUG-2001) National Institutes of Fice, National Cancer Gene Collection (MGC), Cancer Genomics Office, National Cancer Gene Collection (MGC), Room 11A03, Bethesda, MD 20892-2590,
http://www.systemsbiology.org
contact: amadan@systemsbiology.org
Anup Madan, Rachel Dickhoff, Jessica
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Mammalia; 1
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Similar to prostate specific ets
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Rodentia;
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  Fahey,
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eene, Mark Ketteman and Anuradha Madan

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STSEEGWTDGEVDSSCGGQPIHLWQFLKELLLKPHSYGRFIRWLNKEKGLFKIEDSAQ
VARLWGVRKNRPAMNYDKLSRSIRGYKKGIIRKPDISQRLVYQFVHPV*
1575 c 520 g 340 t
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/db_xref="G1:15215055"
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TLDEHSLEQVOSMVVEFYLKDIETACKLLNITADPGDWSPGNVQKWLLWTEHQYRLPP
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427. .1404
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/clone_lib="NCI_CGAP_Co24"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="Mus musculus"
/db_xref="taxon:10090"
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/tissue_type="Colon, normal. 5 m/
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                                                                                                                                                                                                                       CTGAAAGAACTGCTGCTCAAGCCCCACAGCTATGGCCGCTTCATCCGCTGGCTCAACAAG
                                                                                                                                                                                                                                     ctcaaggagttgctactcaagccccacagctatggccgcttcattaggttggctcaacaag
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                                             AAGGGCATCATTCGTAAACCCGACATCTCTCAGCGCCTTGTCTACCAATTTGTGCATCCA
                                                           ATGAAGGAGGACCTCGCCTGGGACCCTTCACTACTGCGCCTCCACCAGCGAGGAGGGC
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During sequence assembly data is compared from overlapping clones. Where differences are found these are annotated as variations together with a note of the overlapping clone name. Note that the variation annotation may not be found in the sequence submission corresponding to the overlapping clone, as we submit sequences with only a small overlap as described above.

This sequence has been finished according to sequence map criteria as follows. An attempt is made to resolve all sequencing problems, such as compressions and repeats, but not necessarily within known annotated repeat sequence elements. Where the sequence is ambiguous, there is an annotation using the 'unsure' feature key. The following abbreviations are used to associate primary accession numbers given in the feature table with their source databases; Em: EMBL; Sw: SWISSPROT' Tr:, TREMBL; Wp:, WORMPEP; Information on the WORMPEP database can be found at http://www.sanger.ac.uk/Projects/C_elegans/wormpep This sequence was generated from part of bacterial clone contigs of human chromosome 6, constructed by the Sanger Centre Chromosome 6 Mapping Group. Further information can be found at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        gccctgcc 1481
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AL157372
AL157372.
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RP11-375E1 is from the library RPGI-11.2 constructed at the Roswell park Cancer Institute by the group of Pieter de Jong. For further details see http://bacpac.med.buffalo.edu/
VECTOR: pBACe3.6
                                                                                                                                                                                                                                                                                                                                                                                                       IMPORTANT: This sequence is not the entire insert of clone RP11-375E1 It may be shorter because we sequence overlapping sections only once, except for a 100 base overlap. The true left end of clone RP11-375E1 is at 1 in this sequence. The true right end of clone RP3-391022 is at 150561 in this sequence. The true right end of clone RP11-302L10 is at 113044 in this
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Submitted (29-NOV-2000) Sanger Centre, Hinxton, Cambridgeshire, CB10 15A, UK. E-mail enquiries: humquery@sanger.ac.uk Clone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HTG
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                                                                                                                                                        /chromosome="6"
/clone="RP11-375E1"
/clone_11b="RPCI-11.2"
/s8887 c 38740 g 37563
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Eutheria;
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/db_xref="taxon:9606"
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Primates;
Score 643; DB 9;
Pred. No. 1.5e-98;
0; Mismatches 5
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                                                    Length 150660;
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Sequence 8 from I
Ax048155
                                                                                                                           Kaufmann, J. and Xin, H.
Expression of ets-domain proteins in Patent: WO 0070092-A 8 23-NOV-2000; Chiron Corporation (US)
                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; V. Mammalia; Eutheria; Primates; Catarrhini; 1 (bases 1 to 1751)
                                                                                                                                                                                        Homo sapiens
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         Similarity
                                                    369
  Conservative
                                                  /note="JKETS promoter"
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/note="TATA box."
521 c 497 g
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/db_xref="taxon:9606"
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         17.4%;
97.7%;
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Patent W00070092.
Score 329.8; DB 6;
Pred. No. 9.8e-46;
0; Mismatches 7;
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AGCAGTTCCGCCAGCGCTCGCCCCTGGGTGGGGATGTGCTGCACGCCCACCTGGACATCT
                                                                                                                                       agcagttccgccagcgctcgcccctgggtggggatgtgctgcacgcccaccttggacatct 1040
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 278)
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (Dases 1 to 278)

1 (Dases 1, 0278)

1 (Dases 2, 0278)

1 (Dases 2, 0278)

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Day,C.H.,
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AX267424
AX267424.1 GI:16516188
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Mammalia; Eutheria;
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Pred. No. 2.2e-34;
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Submitted (17-SEP-2001) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA On Dec 20, 2001 this sequence version replaced gi:16901667.
On Dec 20, 2001 This sequence version replaced gi:16901667.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             * NOTE: Estimated insert size may differ from sequence length

(see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).

* NOTE: This is a 'working draft' sequence. It currently

* consists of 40 contigs. The true order of the pieces

* is not known and their order in this sequence record is

* arbitrary. Gaps between the contigs are represented as

* runs of N, but the exact sizes of the gaps are unknown.

* This record will be updated with the finished sequence
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be preserved.
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/db_xref="taxon:10116"
/clone="CH230-10E14"
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80237: contig
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85983: contig
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ω	1870	98.7	1894	22	AAC83261	Gland-specific Ets
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Ç	1865.2	98.5	3317	22	AAC83266	Gland-specific Ets
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7	1018.6	53.8	1087	21	AAF21828	Human breast and o
æ	738.4	39.0	1704	22	AAH47006	Mouse ETS suppress
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ALIGNMENTS

RESULT AAZ50691

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AAZ50691 standard; cDNA; 1894 BP

Human prostate derived Ets factor cDNA.

23-MAY-2000 AAZ50691;

(first entry)

antibacterial; vulnerary; neuroprotective; antiparkinsonian; nootropic; anabbolic; antiinflammatory; anorectic; hybridisation probe; forensic; tumour marker; diagnosis; treatment; prostate cancer; blood coagulation; autoimmune disorder; haematopoietic; immune/nervous system; stroke; neoplasm; microbial infection; tissue regeneration; heart attack; scarring; food additive; preservative; ss. Prostate Derived Ets factor; PDEF; human; chromosome 6p21.3; cancer; loss of heterozygosity; chromosomal translocation; linkage analysis; cytostatic; cardiant; immunosuppressive; cerebroprotective; fungicide; Key WO200006589-A1 Homo sapiens. Location/Qualifiers 416..1423 /*tag= a /product= "Human prostate derived Ets factor" /note= "Expressed in prostate epithelium"

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                                                                                                                                                                                                                                                                                                                                                                                                                             immunosuppressive, cerebroprotective, fungicide, antibacterial, vulnerary, neuroprotective, antiparkinsonian, nootropic, anabolic, antinfilammatory and anorectic activity, PDEF polynucleotides are usefu in linkage analysis as markers, as hybridisation probes for differential identification of the tissues or cell types and as polymorphic markers for forensic purposes. PDEF is useful as prostate-specific tumour markers for the diagnosis and treatment of prostate cancer. PDEF sequences are useful for treating autoimmune disorders, haematopoietic, blood coagulation, immune and nervous system disorders, hyperproliferative disorders like, neoplasms and microbial infections, heart attacks, stroke, scarring and for tissue regeneration. They are also useful as food additives or preservatives.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present cDNA sequence encodes a human prostate derived Ets (PDEF); 1solated from human prostate carcinoma cDNA library (ATCC No. 203072), PDEF gene expression is highest in tissues androgen sensitivity, with enriched expression in prostate epi cells. PDEF gene is mapped to the human chromosome 6p21.3 region that is associated with loss of heterozygosity and chromosome that is associated with loss of heterozygosity and chromosome form.
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The invention modifying the

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                                    the activity of the gene product. The esf protein or its modified form are useful for inhibiting the growth or proliferation of and/or inducing apoptosis of cancer cells, especially prostate, breast, colon or cervical cancer cells. They are useful for treating cancer in a subject when introduced in to the vicinity of the cancer in a subject. The methods provided are useful for inhibiting the specific degradation of esf protein in a cell; for diagnosing a preneoplastic or neoplastic condition in a human by e.g. PCR and for identifying a compound that inhibits the degradation of esf protein in cancer cells. The present sequence represents a human esf protein encoding cDNA.
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                                                                       CC This invention relates to a method for the detection and determination of CC the metastatic potential of a cell. The method comprises detecting a CC gland-specific Ets transcription factor (GSEF) gene product in a test CC sample. Detection of a GSEF gene product in the test sample in amount CC lower than that in a normal cell, is indicative of a cell with high CC metastatic potential. The method is useful for determining the metastatic potential of a cell, for the diagnosis and prognosis of cancer as well as CC product in test sample. The method may also be used to monitor patients in the producing antibodies, in cancer diagnosis, prognosis, care useful for producing antibodies, in cancer diagnosis, prognosis, care useful for producing antibodies, in cancer diagnosis, prognosis, care useful in gene therapy. GSEF gene product expression levels can be used to conjunction with any tissue in which an alteration in GSEF gene product expression levels is associated with development of a cancer associated phenotype. Cancers, which can be monitored include CC cancers of the prostate, cervix, lung and colon, melanomas, colorectal cadenocarcinomas, Wilm's tumour, retinoblastoma, sarcomas, myosarcomas, CC lung carcinomas, leukaemia, and lymphomas. The GSEF gene is located on the DNA encoding GSEF.
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04-AUG-2001

(first

entry)

Human transcription factor, prost-ets CDNA

antisense Human; transcription vaccine; cancer; pros anscription factor protein;
cancer; prostate; breast; ov
therapy; ribozyme therapy; n; PROST-Ets; c; ovary; benign p y; therapeutic; cytostatic; gene ther
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c; cell proliferation;

CDS Location/Qualifiers 422.1429 /*tag= a /product= "Human PROST-Ets

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Best Local Similarity
Matches 1878; Conserv
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This invention relates to a method for the detection and determination the metastatic potential of a cell. The method comprises detecting a gland-specific Ets transcription factor (GSEF) gene product in a test sample. Detection of a GSEF gene product in the test sample in amount lower than that in a normal cell, is indicative of a cell with high
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cc metastatic potential. The method is useful for determining the metastatic co potential of a cell, for the diagnosis and prognosis of cancer as well as cyrading and staging of cancers by detecting GSEF expression in a complete the method may also be used to monitor patients controlled the prognosis predisposition to develop a particular cancer. GSEF polypeptides are useful for producing antibodies, in cancer diagnosis, prognosis, cyrading, staging and management of breast and prostate tumours, and in conjunction with any tissue in which an alteration in GSEF gene useful in gene therapy. GSEF gene product expression levels can be used conjunction with any tissue in which an alteration in GSEF gene product expression levels is associated with development of a cancer associated phenotype. Cancers, which can be monitored include cancer of the prostate, cervix, lung and colon, melanomas, colorectal cadenocarcinomas, Wilm's tumour, retinoblastoma, sarcomas, myosarcomas, colorectal cancers of the prostate, cervix, lung and colon, melanomas, located on chuman chromosome 6, specifically at 6p21.1-6p21.3. The present sequence corporesents the cDNA encoding GSEF.
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                                           CC The present sequence encodes a human prostate associated Ets protein CC (PRAEP). PRAEP may be used in the prevention, treatment and diagnosis CC of diseases associated with inappropriate PRAEP expression such as CC cell proliferation, immunological, reproductive and developmental CC disorders. The PRAEP polypeptides may be used as antigens in the CC production of antibodies against PRAEP and in assays to identify CC modulators (agonists and antagonists) of PRAEP expression and activity. The anti-PRAEP antibodies and PRAEP antagonists may also be used to CC down regulate PRAEP expression and activity. The anti-PRAEP antibodies CC and antagonists may be used to prevent and treat cell proliferation CC disorders (e.g. cancer, arteriosclerosis and psoriasis), immunological CC disorders (e.g. acquired immunodeficiency syndrome (AIDS), multiple csclerosis and diabetes), reproductive disorders (e.g. polycystic ovary CC syndrome and ectopic pregnancy) and disorders of growth and development CC (e.g. gonadal dysgenesis and achondroplastic dwarfism).
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                                                                                                                                                                                                                                                                                                                                                              CC Sequences AAF21614 - AAF22031 represent DNA sequences encoding human CC proteins AAB58711 - AAB59128. The DNA and protein sequences are CC associated with breast and ovarian cancer. Included in the invention are Sequences AAF22032 - AAF22040 and AAB59129 which are used in the CC isolation and characterisation of the DNA and protein sequences of the CC invention. The breast and ovarian cancer associated DNA, protein, agonist CC or antagonist sequences exhibit cytostatic; immunosuppressive; CC national sequences exhibit cytostatic; immunosuppressive; CC antidiabetic; antiinflammatory; antiviral; antiallergic; hepatotropic; CC antidiabetic; antiinflammatory; antivirer; vulnerary; anticonvulsant; CC antidiabetic; antifungal; antiparastic and cardiant activity. The CC polynucleotide and protein sequences are used in the diagnosis of cancer, CC proteins, agonists and agonists may also be used in the diagnosis, CC proteins, agonists and agonists may also be used in the diagnosis, CC allergies, autoimmune haemolytic anaemia, autoimmune thyroiditis, CC diabetes mellitus, Crohn's disease, multiple sclerosis, rheumatoid CC arthritis and ulcerative colitis; cardiovascular disporders such as CC myocardial ischaemias; wound healing; neurological diseases such as CC cerebral anoxia and epilepsy; and infectious diseases.
                                                                                                                                                                                                                                                                     Query Match 53.8%;
Best Local Similarity 97.0%;
Matches 1055; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New human breast and ovarian cancer associated gene sequences and the polypeptides encoded by these genes, useful in the prevention, treatment and diagnosis of cancer, immune disorders, cardiovascular disorders and neurological diseases -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                autoimmune thyroiditis; diabetes mellitus; Crohn's disease; multiple sclerosis; rheumatoid arthritis; ulcerative colitis; cardiovascular disorder; wound healing; neurological disease;
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                                                                                                                                                The present invention describes a library of polynucleotides comprising CC 1079 nucleotide sequences (given in AAH31145). Also described care: (1) an isolated polynucleotide (I) having at least 90% identity to come of the 1079 sequences; (2) a recombinant host cell containing (I); (CC (3) an isolated polypeptide (II) encoded by (I); (4) an antibody that compressing the polynucleotide (II); (5) a vector comprising (I); and (6) a method compressing the product encoded by (I); (5) a vector comprising the product encoded by (I); (5) of the 1079 sequences given in the specification. The polynucleotides care used to monitor patients having (or susceptible) to cancer to detect consectuable at a gross morphological level. The polynucleotides are useful for monitoring the efficacy of various therapies and preventive consectuations. Polynucleotide probes based on the disclosed sequences care useful for chromosome mapping and detection of transcription levels. The 1079 polynucleotide sequences were derived from a human colon cancer consecutions.
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Matches
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(HYSE-)
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Randazzo F, Kennedy GC, Pot D, Kassam A;
Crkvenjakov R, Dickson M, Drmanac S, Labat
Garcia V, Jones LW, Stache-Crain B;
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antiinflammatory; antithyroid; antiallergic; antibacterial; cardiant; dermatological; neuroprotective; thrombolytic; coagulant; nootropic; vasotropic; antipacoriatic; antianglogenic; gene therapy; inflammation; immune disorder; haematopoietic cell disorder; autoimmune disorder; haematopoietic cell disorder; autoimmune disorder; allergic reaction; graft versus host disease; organ rejection; haemostatic; thrombolytic; cardiovascular disorder; infection; neurological disease; drug screening; ss.
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antidiabetic; antiasthmatic; antirheumatic; antiarthritic; antiviral;
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                                                                                                  Novel isolated nucleic acids comprising sequences encoding useful for treating or diagnosing e.g. cancer - \,
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AAC77607 to AAC78448 encode the human cancer associated proteins given AAB43398 to AAB44239. The proteins can have activities based on tissues and cells the genes are expressed in Example of activities include: cytostatic; proliferative: vulnerary.

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Claim 1; Page 720-721;

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Best Local Similarity
Matches 390; Conser
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                               Transcription factor; gland-specific Ets transcription factor; GSEF; metastatic potential; cancer; tumour; metastasis; breast; prostate; leukaemia; lymphoma; sarcoma; melanoma; chromosome 6p21.1-6p21.3; ds
   Homo sapiens
                                                                                                                        16-MAR-2001
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                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 345; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Detecting metastatic and potential metastatic cancerous cells, useful for diagnosing, prognosing, grading and staging of cancers by detecting gland-specific Ets transcription factor gene product in a biological
  1520
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                   ccagcaggcagccctgggctgggggtaggggactccctacaggcacgcagccctgagacc
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97.78;
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                                                                                                                                                                                                                                          Score 329.8; Db 22,
Pred. No. 3.8e-60;
Pred. No. 3.8e-7;
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14-JUL-1998;
23-SEP-1998;
23-SEP-1999;
15-JAN-1999;
15-JAN-1999;
09-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                The present invention describes isolated polypeptides, comprising an immunogenic portion of a prostate tumour protein (PTP). The polypeptides and polynucleotides encoding them have cytostatic activity and can be used in vaccines and in gene therapy. The polypeptides and polynucleotides encoding them, antigen presenting cells which express the polypeptides, antibodies squainst the polypeptides and vaccines comprising them can be used for inhibiting the development of prostate cancer in a patient. The polypeptides can be used to generate antibodies or anti-idiotypic antibodies for passive immuno therapy. A portion of the polynucleotides encoding the polypeptides can be used as a probe or commodulate the expression of the polypeptides. AAA06241 to AAA06691 and AAV82020 to AAV82020 represent sequences used in the exemplification of
                                               1041
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ggaagtcagcggcctggatgaaagagcggacttcacctggggcgattcactactgtgcct
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                                                                                                                                                                                             Similarity 99.0
53; Conservative
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                                                                                                                                                                                                                                                                                                                    BP;
                                                                                                                                                                                                                                                                                                                                                                    invention.
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RESULT 1
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10-AUG-2000;
29-AUG-2000;
06-SEP-2000;
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10-OCT-2000;
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12-MAY-2000;
13-JUN-2000;
                      polynucleotides, polypeptides, fusion proteins of the polypeptides, antibodies raised against the polypeptides (or antigenic epitopes derived from them) and antigen-presenting cells expressing the polypeptides. The antibodies are useful for detecting the presence of cancer, especially prostate cancer. The polypeptides, polynucleotides and the antigen-presenting cells are useful for stimulating and/or expanding T cells specific for a tumour protein, and for inhibiting the development of cancer especially prostate cancer. Compositions comprising the polynucleotide and/or polypeptide are useful for stimulating an immune response, and for treating cancer. The oligonucleotide is useful for detecting cancer. The present sequence is a prostate specific polynucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAS63826 standard; cDNA;
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                                                                                                                                                                                                                    Claim 1;
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                                                                                                                                                                                                                                            human prostate-specific diagnosis and treatment
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2000US-0568100.

2000US-0570737.

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2000US-0636215.

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2000US-0651236.

2000US-0657279.

2000US-0685166.
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                                                                                                                                                                                         relates to isolated prostate-specific
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                                                                                                                                                                                                                                            polypeptides and polynucleotides useful of cancer, especially prostate cancer -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cytostatic;
                                                                                                                                                                                                                    English.
                                                                                                                                                                                                                                                                                                            , Harlocker SL, Jiang Y, JA, Day CH, Vedvick TS, Hepler WT, Henderson RA;
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Best Local S
Matches 263
The present invention describes polynucleotide sequences (I) which encode prostate-specific proteins (II). (I) and (II) have cytostatic activity, and can be used in vaccine production and gene therapy. (I), (II), antibodies to (II), fusion proteins comprising (II), and isolated T cells prepared using (I) or (II) are used treat cancer in a patient. (I) and the antibodies are also used in the detection of cancer in a patient. The cancer that is diagnosed or treated is particularly prostate cancer. (I) and (II) can be used for monitoring the progression of cancer in a patient. (I) and (II) can also be used to improve diagnostic and therapeutic
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                                                                                                                                                                                                                                                                                                              New polynucleotide encoding a prostate-specific protein, diagnosing, monitoring and treating prostate cancer in a for use in vaccines - \,
                                                                                                                                                                                                                                                                                                                                                                                                               WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Wang
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Kalos
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                                                                                                                                                                                                                                                                      Claim 1;
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                                                                                                                                                                                                                                                                      Page 363;
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, Stolk JA,
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Matches
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                                                                                                                                                                                                                                                                                        Human; prostate cancer; therapy; diagnosis; cat eye syndrome; chromosome 22q11.2; prostate-specific protein; chromosome 1;
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                                               Isolated polypeptide comprising prostate-specific protein, usefu prostate cancer -
                                                                                                                                                                                              09-NOV-2000;
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18-NOV-1999;
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99US-0443686.
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Pred. No. 3e-46;
0; Mismatches
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Claim 31; Page 263; 325pp; English

The present invention describes an at least an immunogenic portion of

isolated polypeptide (P1) comprising a prostate-specific protein, or its

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Matches 263; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           variant. Also described are polynucleotides (N1) encoding (P1). (P1) and (N1) have cytostatic activity and can be used in vaccine production. The polypeptides, nucleic acids and antibodies from the present invention are useful in the diagnosis and therapy of prostate cancer. Prostate specific genes P704P, P712P, P774P, P775P and B305D are located in a genomic region on chromosome 22q11.2 known as the Cat Eye Syndrome region. Prostate specific antigen (PSA) P501S was located on chromosome 1. AAH84671 to AAH85143 and AAG99000 to AAG99077 represent polynucleotide and polypeptide sequences used in the exemplification of the present invention.
                                                                                                                      1221 ttaggtggctcaacaaggagaagg 1244
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99.6%;
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Maximum DB seq length: 2000000000
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

1 3 3 3 4 4 4 4 4 4 1 1 1 1 1 1 1 1 1 1	Result
1894 1894 1870 1870 1870 1870 1868.4 1868.4 1868.4 1868.4 1865.2	Score
100 0 100 0	Query Match 1
1894 1894 1894 1894 1894 1894 1902 1902 1902 1902 1903 1903 1905 1908 11087 11087 1700 470 470 450	Length I
33120554 55 55 55 55 55 55 55 55 55 55 55 55	DB.
PCT-US01-03640-1 US-09-126-945A-1 US-09-567-970-14 US-09-567-970-15 US-09-567-970-15 US-09-166-39-12 US-09-172-373-4734 US-08-4845-995-12 US-09-172-373-4734 US-09-172-373-4734 US-09-172-373-4734 US-09-172-373-4734 US-09-172-373-4734 US-09-172-373-4734 US-09-172-373-4734 US-09-1845-95588 US-09-164-869-277 US-09-644-869-277 US-09-644-869-277 US-09-535-897-20568 US-09-535-897-20568 US-09-535-897-20568 US-09-933-524A-87128 US-09-933-524A-87128	ID
Sequence 1, Appli Sequence 15, Appli Sequence 15, Appli Sequence 17, Appli Sequence 175, Appli Sequence 1637, Appli Sequence 1637, Appli Sequence 3, Appli Sequence 3, Appli Sequence 2057, Appli Sequence 3, Appli Sequence 2058, Appli	Description

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RESULT 1
PCT-USO103640-1
PCT-USO103640-1
Sequence 1, Application PC/TUSO103640
GENERAL INFORMATION: defined and compositions for TITLE OF INVENTION: Methods and compositions for TITLE OF INVENTION: based on ESF
FILE REFERENCE: 9175-015-228
CURRENT APPLICATION NUMBER: PCT/USO1/03640
CURRENT FILING DATE: 2001-02-09
NUMBER OF SED ID NOS: 6
SOFTWARE: Patentin version 3.0
SEQ ID NO 1
LENGTH: 1894
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; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (416)..(1423)
PCT-US01-03640-1
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Matches 1894;
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US-09-933-524-84444
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US-09-240-371-5588
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Pred. No. 0;
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| Sequence 1, Application US/09841963A | GENERAL INFORMATION: APPLICANT: Watson, Dennis K. | APPLICANT: Papas, Takis S. (Deceased) | APPLICANT: Papas, Takis S. (Deceased) | APPLICANT: Papas, Tula C. (Legal Representative) | APPLICANT: Papas, Tula C. (Legal Representative) | TITLE OF INVENTION: based on transcription factor ETS2 | TITLE OF INVENTION: based on transcription factor ETS2 | TITLE OF INVENTION: based on transcription factor ETS2 | CURRENT APPLICATION NUMBER: US/09/841,963A | CURRENT APPLICATION NUMBER: PCT/US99/27805 | PRIOR APPLICATION NUMBER: PCT/US99/27805 | PRIOR APPLICATION NUMBER: PCT/US99/27805 | PRIOR APPLICATION NUMBER: 06/109,850 | PRIOR APPLICATION NUMBER: 06/109,850 | PRIOR APPLICATION NUMBER: 08/109,850 | PRIOR APPLICATION NUMBER: DEGION NUMBER: PATENTAL 1998-11-25 | NUMBER: Patentin version 3.0 | SEQ ID NO 1
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; ORGANISM: Homo sapiens
; FEATURE:
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Sequence 1, Application PC/TUS9917470
GENERAL INFORMATION:
APPLICANT: Human Genome Sciences, Inc.
APPLICANT: Beth Israel Deaconess Medical
APPLICANT: Libermann, Towia A.
APPLICANT: Chitgen, Joerg P.
APPLICANT: Kunsch, Charles A.
APPLICANT: Endress, Gregory A.
APPLICANT: Rosen, Craig A.
APPLICANT: Rosen, Craig A.
APPLICANT: Rosen, Craig A.
APPLICANT: Rosen, Craig A.
CURRENT APPLICATION NUMBER: PCT/US99/174:
CURRENT APPLICATION NUMBER: PCT/US99/174:
CURRENT FILING DATE: 1998-07-31
NUMBER OF SEG ID NOS: 15
SOFTWARE: Patentin Ver. 2.0
                                                    ; LENGTH: 1894
; TYPE: DNA
; ORGANISM: HOMO :
PCT-US99-17470-1
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PCT-US99-17470-1
Sequence 1, Application PC/TUS9917470
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US-09-126-945-1
Sequence 1, Application US/09126945
GENERAL INFORMATION:
APPLICANT: Libermann, Towia A.
APPLICANT: Libermann, Towia A.
APPLICANT: Chetten, Joerg P.
APPLICANT: Endress, Gregory A.
APPLICANT: Rosen, Craig A.
TITLE OF INVENTION: Prostate Derived Ets
FILE REFERENCE: 1488.1090000
CURRENT APPLICATION NUMBER: US/09/126,94
CURRENT FILING DATE: 1998-07-31
NUMBER OF SEQ ID NOS: 15
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 1
LENGTH: 1894
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Libermann, Towia A. Oettgen, Joerg P. Kunsch, Charles A. Endress, Gregory A.
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TITLE OF INVENTION: Prostate Deri
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APPLICANT: Libermann, Towia A.
APPLICANT: Oettygen, Joerg P.
APPLICANT: Cottygen, Charles A.
APPLICANT: Endress, Gregory A.
APPLICANT: Endress, Gregory A.
APPLICANT: ROSEN, Craig A.
TITLE OF INVENTION: PROSTATE Derived Et
FILE REFERENCE: 1488.1090000
CURRENT APPLICATION NUMBER: US/09/126,94
CURRENT FILING DATE: 198-07-31
NUMBER OF SEQ ID NOS: 15
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 1
LENGTH: 1894
TYPE: DNA
ORGANISM: HOME Sapiens
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; GENERAL INFORMATION:
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SOFTWARE: FASTSEQ for Wind
SEQ ID NO 1
LENGTH: 1894
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
FEATURE:
NAME/KEY: CDS
LOCATION: (416)...(1423)
OTHER INFORMATION: Human J
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Best Local Similarity 99.2%;
Matches 1879; Conservative
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CURRENT FILING DATE: 2000-05-12
PRIOR APPLICATION NUMBER: 60/134,112
PRIOR FILING DATE: 1999-05-14
NUMBER OF SEO ID NOS: 13
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APPLICANT: Xin, Hong
APPLICANT: Harrowe, Greg
TITLE OF INVENTION: EXPRESSION
TITLE OF INVENTION: CANCER
FILE REFERENCE: 2300-1556
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                                                                                           FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION UMBER: 08/845,99
APPLICATION NUMBER: 08/845,99
APPLICATION NUMBER: 35,441
REGISTRATION NUMBER: 35,441
REFERENCE/DOCKET NUMBER: 6085
TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/933-1729
TELEPAX: 847/938-2623
                                                                     NFORMATION FOR SEQ ID NO:
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APPLICANT:
APPLICANT:
APPLICANT:
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OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/064
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MEDIUM TYPE: Diskette
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STREET: 100 Abbott
CITY: Abbott Park
STATE: IL
                                                         SEQUENCE CHARACTERISTICS:
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TYPE: nuclear sin
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JOHN C. RUSSELL
STEPHEN D. STROUP
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PAULA N. FRIEDMAN
MICHAEL R. KLASS
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RESULT 9
US-09-064-645-15
: Sequence 15, Application US/09064645
: Sequence 15, Application US/09064645
: GENERAL INFORMATION:
APPLICANT: MAURICE COHEN
: APPLICANT: MAURICE COHEN
: APPLICANT: TRACEY L. COLDITTS
APPLICANT: MICHAEL R. KLASS
APPLICANT: LISA ROBERTS-RAPP
: APPLICANT: LISA ROBERTS-RAPP
: APPLICANT: JOHN C. RUSSELL BILLING-MEDEL

540

600

660

780

787

667 720 727

840

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COUNTRY: LA

ZIP: 60064-3500

ZOUNTRY: USA
ZIP: 60064-3500

COMPUTER READABLE FORM:
MEDIOM TYPE: Diskette
COMPUTER: IBM. COMPATIBLE
OPERATING SYSTEM: DOS
SOFTWARE: FRSYSEQ: for Windows Ver
CURRENT APPLICATION NUMBER: US/09/064,64
FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION NUMBER: 08/845,995
FILING DATE: 25-APR-1997
ATTORNEY/AGENT INFORMATION:
NAME: Becker, Cheryl L.
REGISTRATION NUMBER: 6085:
REFERENCE/DOCKET NUMBER: 6085:
TELECOMMUNICATION INFORMATION:
TELECHONE: 847/935-1729
TELEEX:
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
LENGTH: 1902 base pairs
TYPE: nucleic acid
STRANDEDNESS: Single
TOPOLOGY: linear
US-09-064-645-15
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STREET: 100 Abbott
CITY: Abbott Park
STATE: IL
COUNTRY: USA
ZIP: 60064-3500
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                                                                                                                                                                                                                                                                                                                        Conservative
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REAGENTS AND MI
FOR DETECTING I
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Sequence 14, Application US/09567970

GENERAL INFORMATION:
APPLICANT: Abbott Laboratories
APPLICANT: Billing Medel, Patricia A.
APPLICANT: Colpitts, Tracey L.
APPLICANT: Criedman, Paula N.
APPLICANT: Friedman, Paula N.
APPLICANT: Russell, John C.
APPLICANT: Russell, John C.
APPLICANT: Stroupe, Stephen D.
TITLE OF INVENTION: REAGENTS AND METHODS USITITE OF INVENTION: DETECTING DISEASES OF TIFLE OF INVENTION: DETECTING DISEASES OF TIFLE OF APPLICATION NUMBER: US/09/567,970
CURRENT APPLICATION NUMBER: US 09/064,645
PRIOR FILING DATE: 1998-04-22
PRIOR FILING DATE: 1997-04-25
NUMBER OF SEQ ID NOS: 37
SCOTUMARE: FastSEQ for Windows Version 4.0
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US-09-
                                                                                                                                                                                                                   SEQ ID NO 14
LENGTH: 1902
TYPE: DNA
ORGANISM: HOMO:
S-09-567-970-14
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Best Local Similarity
Matches 1878; Conserv
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Sequence 15; Applicant INFORMAL

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FILE REFERENCE

CURRENT FILING DA

PRIOR APPLICAT

PRIOR APPLICAT
                                                                               APPLICANT: Abbott Laboratories
APPLICANT: Billing-Medel, Patricia A.
APPLICANT: Colpitts, Tracey L.
APPLICANT: Colpitts, Tracey L.
APPLICANT: Colpitts, Tracey L.
APPLICANT: Klass, Michael R.
APPLICANT: Klass, Michael R.
APPLICANT: Roberts-Rapp, Lisa
APPLICANT: Russell, John C.
APPLICANT: Stroupe, Stephen D.
TITLE OF INVENTION: DETECTING DISEASES OF THITLE OF INVENTION: DETECTING DISEASES OF THIFILE OF INVENTION NUMBER: US/09/567,970
CURRENT APPLICATION NUMBER: US/09/567,970
CURRENT APPLICATION NUMBER: US 09/064,645
PRIOR FILING DATE: 1998-04-22
PRIOR APPLICATION NUMBER: US 08/845,995
PRIOR APPLICATION NUMBER: US 08/845,995
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LENGTH: 1902
TYPE: DNA
ORGANISM: Homo :
3-09-567-970-15
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RESULT 12
US-09-718-159-1
Sequence 1, Application
GENERAL INFORMATION:
APPLICANT: Harkins, APPLICANT: Liho, Rici
APPLICANT: Luke, Maj
APPLICANT: Monteclai
APPLICANT: Monteclai
APPLICANT: Parkes, I
APPLICANT: Parkes, I
APPLICANT: Steinbree
APPLICANT: Van Heuit
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Luke, May
Monteclaro, Felipe
Parkes, Deborah
Parry, Gordon
Steinbrecher, Renate
Van Heuit, Pamela Toy
                                                                             Application
                                                   Lin, Richard
                                                            Harkins, Richard
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; APPLICANT: Xuan, Jian-ai
; TITLE OF INVENTION: DNA |
; FILE REFERENCE: 51790AUS;
; CURRENT APPLICATION NUMBE;
; CURRENT FILING DATE: 200(
; PRIOR APPLICATION NUMBER:
; PRIOR FILING DATE: 1999-1
; NUMBER OF SEQ ID NOS: 11
; SOFTWARE: Patentin Ver. 2
; SEQ ID NO 1
; LENGTH: 1918
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: CDS
; LCCATION: (422)..(1429)
US-09-718-159-1
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TITLE OF INVENTION: DNA Encoding a Novel ET; FILE REFERENCE: 51790AUSM1
; CURRENT APPLICATION UNMBER: US/09/718,159
; CURRENT FILING DATE: 2000-11-21
; PRIOR APPLICATION NUMBER: 60/168,182
; PRIOR FILING DATE: 1999-11-30
; NUMBER OF SEQ ID NOS: 11
; SOFTMARE: Patentin Ver. 2.0
; SEQ ID NO 1
; LENGTH: 1918
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LENGTH: 3317

TYPE: DNA

ORGANISM: Homo sapiens
FEATURE:
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OTHER INFORMATION: GSEF-encoding se
NAME/KEY: promoter
LOCATION: (1)...(1380)
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APPLICANT: Harrowe, GI
APPLICANT: Harrowe, TITLE OF INVENTION: EX
TITLE OF INVENTION: CA
FILE REFERENCE: 2300-1
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Best Local Similarity
Matches 1876; Conser
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CURRENT FILING DATE: 2000-05-12
PRIOR APPLICATION NUMBER: 60/134,112
PRIOR FILING DATE: 1999-05-14
NUMBER OF SEQ ID NOS: 13
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RESULT 14

US-09-866-356-2

Sequence 2, Application US/09866356

GENERAL INFORMATION:

APPLICANT: Bandman, Olga

CORILLY, Neil C.

Guegler, Karl J.

ITILE OF INVENTION: PROSTATE-AS

NUMBER OF SEQUENCES: 4

CORRESPONDENCE ADDRESS:

ADDRESSE: Incyte Pharmace

STREET: 3174 Porter Dr.

CITY: Palo Alto

STATE: CA

COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette

COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette

COMPUTER: IBM Compatible

OPERATING SYSTEM: DS

SOFTWARE: FastSOD for Wind

CURRENT APPLICATION NUMBER: US/09/

FILING DATE: 1998-04-03

ATTORNEY/ACENT INFORMATION:

REGISTRATION NUMBER: 09/055

TELEPHONE: 650-85-0555

TELEPHONE: 650-85-0555

TELEFAX: CUNKNOWN>

INFORMATION FOR SEQ ID NO: 2:

SEQUENCE CHARACTERISTICS:

LENGTH: 1905 base pairs

TYPE: NUCLEIC acid

STRANDEDURSS: single

TODOLOGY: Linear

INTERNATOR DESCRIPTION

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FILING DATE: 1998-04-03
ATTORNEY/AGENT INFORMATION:
NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-0
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-855-055
TELEPAX: 650-845-4166
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                                                                       LIBRARY:
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                        DESCRIPTION:
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SYSTEM: DOS
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RESULT 15
US-60-209-009-74
US-60-209-009-74
Sequence 74, Application US
GENERAL INFORMATION:
APPLICANT: Faris, Mary
APPLICANT: Pearson, Cece,
TITLE OF INVENTION:
FILE REFERENCE: PA-0027
CURRENT APPLICATION NUMBE:
CURRENT FILING DATE: 200;
NUMBER OF SEQ ID NOS: 50

US/60209009

APPLICANT: Faris, Mary
APPLICANT: Pearson, Cecelia I.
APPLICANT: Pearson, Cecelia I.
TITLE OF INVENTION: GENES EXPRESSED:
FILE REFERENCE: PA-0027 P
CURRENT APPLICATION NUMBER: US/60/209,009
CURRENT FILING DATE: 2000-06-01
NUMBER OF SEQ ID NOS: 501

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ORGANISM: HOMO sapiens
FEATURE:
NAME/KEY: misc_feature
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Sequence 10 from Patent WO0070092.
AX048157
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                                                                                                                                                                         Harkins,R., Lin,R.J., Luke,M., Monteclaro,F., Steinbrecher,R., van Heult,P.T. and Xuan,J.A. Dna encoding a novel prost-ets polypeptide Patent: WO 0142472-A 5 14-JUN-2001; SCHERING AKTIENGESELLSCHAFT (DE)
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Sequence 5 from Patent WO0142472.
AX175520
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1 (bases 1 to 23)
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/db_xref="taxon:32630"
/note="GSEF-specific primer"
/note="13 g 1 t
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/db_xref="taxon:32630"
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M synthetic construct
artificial sequence.

1 (bases 1 to 23)
Harkins,R., Lin,R.J., Luke,M., Monteclaro,F., Pa
Steinbrecher,R., van Heult,P.T. and Xuan,J.A.
Dna encoding a novel prost-ets polypeptide
Patent: WO 0142472-A 10 14-JUN-2001;
SCHERING AKTIENGESELLSCHAFT (DE)
TGACTCGACAAAGGCCACAGGCA 23
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Sequence 10 from Patent WO0142472.
AXI75525
AXI75525.1 GI:14598861
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Sequence 9 from Patent WO0142472.
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Harkins,R., Lin,R.J., Luke,M., Monteclaro,F.,
Steinbrecher,R., van Heuit,P.T. and Xuan,J.A.
Dna encoding a novel prost-ets polypeptide
Patent: WO 0142472-A 914-JUN-2001;
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/db_xref="taxon:32630"
/note="antisense"
8 c 5 g 6 t
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/note="antisense"
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Nucleic acids containing single nucleotide
methods of use thereof
Patent: WO 0151670-A 618 19-JUL-2001;
Curagen Corporation (US)
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Sequence 618 from Patent WO0151670.
AX199688 AX199688.1 GI:15390129
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Sequence 11 from Patent WO0142472.
AX175526
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Eukaryota; M
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Similarity 66.0%;
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/db_xref="taxon:9606"
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/db_xref="taxon:32630"
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Pred. No. 4.8e+06;
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                                                                                                                                                                                                                                                                                                              JOURNAL
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                                                                                                                              59
                                                                                                     1 CCAGTGCTCCAAGCCYGGTGTCATCTTCCT 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 GGCTGGGCGGAGGAAACTCCCTTCAAGGACATCGTCCGGGGACTG 45
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                                                                                                                 ccagtcctccaagcctgctgccagctccct 88
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MUSJHCD6
Mouse JH-Cdelta locus
M64562
M64562.1 GI:198478
                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Mammalia; Eutheria; Primates; Catarrhini (Dases 1 to 31)
Cargill, M., Ireland, J.S. and Lander, E.S.
                                                                                                                                                                                                                                                                                                                                                                                                                            AX249190
Sequence 1269 from Patent
AX249190
AX249190.1 GI:15863813
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            artificial sequence.

1 (bases 1 to 48)
Blatt.L. Mcswiggen,J. and Chowrira,B.M.
Method and reagent for the modulation an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 7387 from Patent AX221945 AX221945.1 GI:15549669
                                                                                                                                                                                                                                                                                 Human single nucleotide polymorphisms
Patent: WO 0166800-A 1269 13-SEP-2001;
WHITEHEAD INSTITUTE FOR BIOMEDICAL RESEARCH
LOCATION/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RIBOZYME PHARMACEUTICALS, INC. (US); Blatt, McSwiggen, James (US); Chowrira, Bharat M. (Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   nogo gene expression Patent: WO 0159103-A 7387 16-AUG-2001;
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ilarity 80.08;
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                                                                                                                                                                                                                                  /organism="Homo sapiens"
/db_xref="taxon:9606"
11 c 6 g 8
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/db_xxef="taxon:32630"
/note="Nucleic Acid"
13 c 18 g 7 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.2%;
                                                                                                                                                     Score 21.6; DB 6;
Pred. No. 8.6e+06;
1; Mismatches 5
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                         49 bp sequence.
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Pred. No. 5.3e+06;
0; Mismatches 14
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WO0159103
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W00166800
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HSTCK6X45
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A05116
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Matches 26
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                                                                                                                   1 CTCGGGCCCTGCAGTTCCTCAGCAGTCTTCA 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CCAAGATGGTGGGCTGGGACAGAGCAGAGCTTGCTGTGGCCCACACC 3
HSTCK6X45
H.sapiens mRNA (clone K6-45).
269575
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1 (bases 1 to 49)
Owens, J.D. Jr., Finkelman, F.D., Mountz, J.D. and Mushinski, J.F. Nonhomologous recombination at sites within the mouse JH-Cdelta locus accompanies Cmu deletion and switch to immunoglobulin D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A05116 45 bp DNA 1i
Oligonucleotide U7 for bovine growth hormone.
A05116
A05116.1 GI:345003
                                                                                                                                                                            1.1%;
L Similarity 76.5%;
26; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                synthetic construct synthetic construct
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                             Patent: WO 8805078-A 13 14-JUL-1988;
                                                                                                                                                                                                                                                                                                                                                         SYNTHETIC GENE
                                                                                                                                                                                                                                                                                                                                                                                                                   artificial sequence.
                                                                                                                                                                                                                                                                                                                                                                        METHOD FOR THE PRODUCTION OF BOVINE GROWTH HORMONE USING
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/db_xref="taxon:10990"
/cell_tine="KyD8"
/cell_type="Hybridoma"
a 17 c 14 g 11 t
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/db_xref="taxon:32630"
17 c 10 g 10 t
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                                                                                                                                                                            Score 21.2; DB 6;
Pred. No. 9.1e+06;
0; Mismatches 8
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Pred. No. 8.3e+06;
0; Mismatches 16
                           48 bp mRNA lir
receptor beta chain
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                                            PRI 05-FEB-1997
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AUTHORS
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TITLE
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JOURNAL
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Best Local
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                                  AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The T-cell receptor beta chain CDR3 region of BV8S1/BJ1S5 transcripts in type 1 diabetes Immunogenetics 45 (2), 87-96 (1996)
                                                                                                                                                                                                                                                                                                                                                                                                                                                29;
                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 48)
Naserke,H.E., Durinovic-Bello,I., Seidel,D. and Ziegler,A.G. The T-cell receptor beta chain CDR3 region of BV8S1/BJ1S5 transcripts in type 1 diabetes
Immunogenetics 45 (2), 87-96 (1996)
                                                                                                                                                                                                                                                                                        H.sapiens
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Naserke, H.E., Durinovic-Bello, I.,
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Direct Submission Submitted (02-FEB-1996) Naserke H.E.,
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Mammalia; Eutheria;
                                  Naserke, H.E.
                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                        diversity region; joining region; T cell receptor; variable region
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                                                                 97113296
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                                                 (bases 1 to 48)
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/db_xref="taxon:9606"
/tissue_type="peripheral blood"
/cell_type="lymphocytes"
/clone="K6-45"
/clone="K6-45"
/note="HLA type: HLA-A31, 1; Cw3, 3; B6
0401; DRB3*0202, -; DQA1*0103, 0301; DQ
DPA1*, 01; DPB1*0301, 0201"
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29. .>48
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29. .48
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12. .28
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13 c 16 g
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for T cell receptor
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Primates;
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    Diabetes Research Institute
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DQB1*0603, 0302;
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                                                                               artificial sequence.

artificial sequence.

1 (bases 1 to 50)

Leadlay, P. F., Staunton, J. and Oliynyk, M.C.

Polyketides and their synthesis

Patent: WO 0168867-A 51 20-SEP-2001;

Biotica Technology Limited (GB)
                                                                                                                                                                                                                                                                                                                                                                                        Sequence 51 from Patent W00168867.
AX175518 21 bp
Sequence 3 from Patent WO0142472.
AX175518
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/tissue_type="peripheral blood"
/ceil_type="lymphocytes"
/clone="K6-45"
/note="ur"
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29. .48
                                                                                                                                                                                                                         /organism="synthetic construct"
/db_xref="taxon:32630"
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12. .2
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AUTHORS
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                                                                       1 AGTGACTCGACAAAGGCCACA 21
                                                                                                                                                                                                                                                                                                artificial sequence.

1 (bases 1 to 21)

Harkins,R., Lin,R.J., Luke,M., Monteclaro,F., Parkes,D., Steinbrecher,R., van Heuit,P.T. and Xuan,J.A.

Dna encoding a novel prost-ets polypeptide

Patent: WO 0142472-A 3 14-JUN-2001;

SCHERING ANTIENGESELLSCHAFT (DE)
                                                                                                                                                                                                                                                                                                                                                                                                      synthetic construct.
synthetic construct
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/db_xref="taxon:32630"
/note="primer"
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100.0%; Pred. No.
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1.2e+07;
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Maximum Match 100%
Listing first 45 summaries
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2: /SIDS5/gcgdata/geneseq/geneseqn-emb1/NA1981.DAT:*
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10425.224 Million cell updates/sec
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SIDS5/gcgdata/geneseq/geneseqn-emb1/NA1985.DAT:

SIDS5/gcgdata/geneseq/geneseqn-emb1/NA1986.DAT:

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                                /SIDS5/gcgdata/geneseq/geneseqn-emb1/NA2000.DAT:*
/SIDS5/gcgdata/geneseq/geneseqn-emb1/NA2001A.DAT:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Query Description Score Match Length DB ID Description January Description AACB3265 GSEF specific PCR GSEF specif	vρœ	76	ი 5	4	c 3	2	c 1	Result
ID 2 AAC83265 2 AAC83264 2 AAS5839 1 AAZ65828 2 AAL07340 2 AAD07341 2 AAD07342 2 AAD07343	23 23	23 23	23.6	23.6	24.6	31	34	Score
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Description GSEF specific PC GSEF specific PC Antisense oligon Human map-relate Human SNP oligon Human transcript Antisense oligo Antisense oligo Antisense oligo	342 343	340 341	612	828	839	264	265	1 † 1 1 † †
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Detecting metastatic and potential metastatic cancerous cells, useful

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AAZ65894 ABK07085 ABK07086	ABK06870 AAI30781 AAC81266	AAX59472 AAX59473 ABL01057	AAT40111 AAV39429 AAV39430	AAL34241 AAA04697 ABK06858 ABK06869 ABA10736	AAT80517 AAT80517 AAF60322 AAQ11438 AAZ23808 AAZ23808 AAL23118 AAL23808 AAL23808	AAF74533 AAC64157 AAC64158	AAD19606 AAD19607 AAF74533	AAL31921 AAH89837 ABK07387 AAL28314 AAL34308
Human map-related Human NOGO Amberzy Human NOGO Amberzy	Sing mous	Primer used in con Primer used in con Human SNP involvin	220	Human SNP oligonuc Polymorphic fragme Human NOGO Amberzy Human NOGO Amberzy Tail adaptor oligo	Hepatoma AS-30D Ty Human SNP oligonuc PDGF-B receptor bi Probe #4 complemen Human Kv6.2 DNA fr Human SNP oligonuc Human transcriptio FIV gene cloning r	MKK3	enic	Human SNP oligonuc Human coding segu Human NGO Amberzy Human SNP oligonuc Human SNP oligonuc

ALIGNMENTS

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RESULT 1
AAC83265/c
ID AAC83265;
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AC83265;
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AC AAC83265;
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Tenascription factor; gland-spec
KW metastatic potential; cancer; tu
Leukaemia; lymphoma; sarcoma; me
KW PCR primer; ss.
XX
PN W0200070092-A1.
XX
PP 12-MAY-2000; 2000WO-US13173.
XX
PF 12-MAY-1999; 99US-0134112.
XX
PA (CHIR ) CHIRON CORP.
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RESULT
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                                                                                                                                                                                                                                                                                      Transcription factor; gland-specific Ets transcription factor; GSEF; metastatic potential; cancer; tumour; metastasis; breast; prostate; leukaemia; lymphoma; sarcoma; melanoma; chromosome 6p21.1-6p21.3;
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               WPI; 2001-041019/05
                                                                                                                                           12-MAY-2000; 2000WO-US13173
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Sequence 31

BP; 8 A; 9 C; 13 G; 1 T; 0 other;

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CC the metastatic potential of a cell. The method comprises detecting a completation of a GSEF gene product in the test sample. Detection of a GSEF gene product in the test sample in amount comments of a cell. Is indicative of a cell with high comments than that in a normal cell, is indicative of a cell with high compression than that in a normal cell, is indicative of a cell with high competential of a cell, for the diagnosis and prognosis of cancer as well as completed in the method is useful for determining the metastatic potential of a cell, for the diagnosis and prognosis of cancer as well as comparing and staging of cancers by detecting GSEF expression in a comparing and staging of cancers by detecting GSEF expression in a comparing a predisposition to develop a particular cancer. GSEF polypeptides care useful for producing antibodies, in cancer diagnosis, prognosis, comparing and management of breast and prostate tumours, and in conjunction with any tissue in which an alteration in GSEF gene concerts of the prostate, cervix, lung and colon, melanomas, colorectal cancers of the prostate, cervix, lung and colon, melanomas, colorectal cancers of the prostate, cervix, lung and colon, melanomas, colorectal colores and prostate and prostate on the prostate on the prostate of primer specific for the human GSEF DNA sequence. The corporate is used for the chromosamal mapping of the GSEF gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Detecting metastatic and potential metastatic cancerous cells, useful for diagnosing, prognosing, grading and staging of cancers by detecting gland-specific Ets transcription factor gene product in a biological sample from a cell -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Page 56;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               relates to a method for the detection and determination potential of a cell. The method comprises detecting a Ets transcription factor (GSEF) gene product in a test
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          95pp;
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RESULT AAX35839/ В δã Query Match Best Local S Matches 31 LT 3 5839/c AAX35839 1552 cagggagggcaaccaactgccccaggggga 1582 WPI; 1999-276982/23. 25-SEP-1998; 01-APR-1999 WO9915643-A2 Synthetic ACE; hypertension; ss Antisense oligonucleotide; mammalian; angiotensin converting enzyme; Antisense oligonucleotide for mammalian ACE mRNA 14-JUL-1999 25-SEP-1997; (UYFL) UNIV 1 cagggaggggaaccaactgccccaggggga 31 Ď Similarity standard; DNA; 42 Conservative FLORIDA (first entry) 97us-0059661 98WO-US20121 100.0%; M 1.6%; Score 31; pred. No. 0; Mismatches DB 1.2e+03; 22; Length 31; Indels 0; Gaps

New antisense oligonucleotide compositions

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Best Local S
Matches 30
invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAX59579 to AAX77440 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human genome; biallelic marker; high density disequilibrium map; genomic map; haplotype; phenotype; potition; biase; genotyping; haplotyping; hybridisation; identification; characterisation; diagnosis; single nucleotide polymorphism; SNP; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      angiotensin converting enzyme (ACE) mRNA. The antisense oligonucleotides can be used for treating or preventing hypertension. The antisense oligonucleotides can also be labeled and used for visualising ACE mRNA in a cell, and producing transgenic animals.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human map-related biallelic marker
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                                                                                                                                                                                                                                                           21-APR-1998;
23-NOV-1998;
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                                                                                                 AAZ65654 to AAZ69578 represent human biallelic markers from the present
                                                                                                                          Claim
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                                                                                                                                                                                                                                                                                               21-APR-1999;
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                                                                                                                                                            biallelic markers
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                                                                                                                                                                                                            Blumenfeld M,
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ilarity 76.9%;
Conservative
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98US-0109732.
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the development of
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Best Local
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and 3367,
The present invention relates to oligonucleotides encoding polymorphic variants of proteins related to amylases, amyloid proteins, angiopoietin, apoptosis related proteins, cadherin, cyclin, polymerase, oncogenes, histones, kinases, colony stimulating factors, complement related proteins, cytochromes, kinesins, cytokines, interferons, interleukins, G-protein coupled receptors and thioesterases. The present sequence is one such oligonucleotide. The oligonucleotides and the peptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of the proteins listed above. Disorders that may be prevented, diagnosed and/or treated include multifactorial diseases with a genetic component, such as autoimmune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Immunosuppressive; immunostimulatory; antiinflammatory; cytostatic; neuroprotective; antimicrobial; gene therapy; vaccine; amylase; cancer; amyloid protein; angiopoietin; apoptosis related protein; cadherin; cyclin; polymerase; oncogene; histone; kinase; colony stimulating facto complement related protein; cytochrome; kinesin; cytokine; interferon; interleukin; G-protein coupled receptor; thioesterase; inflammation; multifactorial disease; autoimmune disease; infection; nervous system disease; ss.
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                                                                                                                                                                                                                        Polymorphic nucleic acids encoding e.g. amylases, oncogenes and histones, useful for diagnosing and cancer. autoimmune diseases and infections -
                                                                                                                                                                                                                                                                                 WPI;
                                                                                                                                                                                                                                                                                                              Shimkets RA,
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27-DEC-2000; 2000US-0173419.
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                                                                                                                                                                                         Claim
                                                                                                                                                                                                                     cancer, autoimmune diseases
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                                                                                                                                                                                       Page 3652; 4143pp;
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Pred. No. 4.8e+04;
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Best Local S
Matches 32
The invention relates to human transcription factor protein, designated as PROST-Ets and its corresponding cDNA molecule. PROST-Ets proteins and nucleic acids are useful for treating a disease-state in human patient, such as prostate, breast and ovarian cancer which is associated with inappropriate expression of PROST-Ets and where the patient is in need of decreased levels or activity of PROST-Ets. PROST-Ets is useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              340
                                                                                                                            New transcription factor polypeptides, designated polypeptides, useful for diagnostic, research and
                                                                                                                                                                                                                                                     30-NOV-1999; 99US-0168182
21-NOV-2000; 2000US-0718159
                                                                                                                                                                                                                                                                                                                                                                                                          modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; transcription factor protein; PROST-Ets; cytostatic; gene therapy;
vaccine; cancer; prostate; breast; ovary; benign prostatic hyperplasia;
antisense therapy; ribozyme therapy; therapeutic; cell proliferation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          diseases (e.g. rheumatoid arthritis, multiple sclerosis, diabetes, systemic lupus erythromatosus and Grave's disease), inflammation, cancer (e.g. cancers of the bladder, brain, breast, colon and kidney, leukaemia), diseases of the nervous system and an infection of pathogenic
                                                                                         Example 2;
                                                                                                                                                                                           Steinbrecher
                                                                                                                                                                                                       Harkins
                                                                                                                                                                                                                                                                                            29-NOV-2000; 2000WO-US42295
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32; Conser
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                                                                                                                                                                                                                                 SCHERING
                                                                                                                                                                                         Lin RJ,
er R, Va
                                                                                        Page 48; 74pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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23
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                                                                                                                                                                                                                                                                                                                                                                                                                      Cytosine"
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21-NOV-2000;
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2000US-0718159
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therapeutic
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The invention relates to human transcription factor protein, designated as PROST-Ets and its corresponding cDNA molecule. PROST-Ets proteins and nucleic acids are useful for treating a disease-state in human patient, such as prostate, breast and ovarian cancer which is associated with inappropriate expression of PROST-Ets and where the patient is in need of decreased levels or activity of PROST-Ets PROST-Ets is useful for diagnosing and treating diseases of cell proliferation, such as prostate cancer and benign prostatic hyperplasia. Prost-ets nucleic acids are used as DNA probes, as target for antisense and ribozyme therapy, as template for producing antisense polynucleotides, and as diagnostic reagent for research, biological, clinical and therapeutic purposes. Prost-ets DNA is

Claim 36; Page 56; 74pp;

English.

research, biological, clinical and therapeutic purposes. Prost-ets DN also used in gene therapy. PROST-Ets protein is useful for generating antibodies, and for stimulating immune response in cells containing

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                       The invention relates to human transcription factor protein, designated CC as PROST-Ets and its corresponding CDNA molecule. PROST-Ets proteins and CC nucleic acids are useful for treating a disease-state in human patient, CC such as prostate, breast and ovarian cancer which is associated with C inappropriate expression of PROST-Ets and where the patient is in need of CC decreased levels or activity of PROST-Ets. PROST-Ets is useful for CC cancer and benign prostatic hyperplasia. Prost-ets nucleic acids are used CC cancer and benign prostatic hyperplasia. Prost-ets nucleic acids are used CC as DNA probes, as target for antisense and ribozyme therapy, as template CC research, biological, clinical and therapeutic purposes. Prost-ets DNA is CC also used in gene therapy. PROST-Ets protein is useful for generating CC antibodies, and for stimulating immune response in cells containing CR ROST-Ets proteins. The present DNA sequence is antisense oligonucleotide which is used for evaluating the role of human transcription factor, the selected profession in cells proliferation to reduce Prost-ets mRNA expression in
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Matches 17; Conserv
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Steinbrecher
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21-NOV-2000; 2000US-0718159
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Pred. No. 5.6e+04
                cell lines
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                        diagnosing and treating diseases of cell proliferation, such as prostate cancer and benign prostatic hyperplasia. Prost-ets nucleic acids are used as DNA probes, as target for antisense and ribozyme therapy, as template for producing antisense polynucleotides, and as diagnostic reagent for research, biological, clinical and therapeutic purposes. Prost-ets DNA is also used in gene therapy. PROST-Ets protein is useful for generating antibodies, and for stimulating immune response in cells containing antibodies, and for stimulating immune response in cells containing PROST-Ets proteins. The present DNA sequence is antisense oligonucleotide which is used for evaluating the role of human transcription factor, PROST-Ets in cell proliferation to reduce Prost-ets mRNA expression in the selected prostate tumour cell lines.
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21-NOV-2000; 2000US-0718159.
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Pred. No. 5.6e+04;
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Query Match Best Local Similarity

1.2%;

Score Pred.

23; No.

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Length

Sequence

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ALISIPZI/c
ID AALISIPZI standard; D
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AC AALISIPZI;
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DE Human SNP oligonucle
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KW Immunosuppressive; ant
KW neuroprotective; and
KW nervous system disea
XX
Homo sapiens.
XX
WO200147944-A2.
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HOMO Sapiens.
XX
WO200147944-A2.
XX
Z8-DEC-2000; 2000WO-
XX
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Z8-DEC-1999; 99US-
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XY
MPI; 2001-465210/50.
XX
PO1ymorphic nucleic
PT oncogenes and histon
PT cancer, autoimmune d
XX
Claim 1; Page 2862;
XX
Yhe present inventio
CC variants of proteins
CC variants of proteins
CC variants of proteins
CC variants of proteins
CC core such oligonucleo
CC proteins, cytochrome
CC proteins, cytochrom
CC proteins, cytochrome
CC proteins, cytochrome
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                                                                                                                                                                                           The present invention relates to oligonucleotides encoding polymorphic CC variants of proteins related to amylases, amyloid proteins, anglopoletin, apoptosis related proteins, cadherin, cyclin, polymerase, oncogenes, cC apoptosis related proteins, cadherin, cyclin, polymerase, oncogenes, cC instenes, kinases, colony stimulating factors, complement related CC proteins, cytochromes, kinesins, cytokines, interferons, interleukins, CC G-protein coupled receptors and thioesterases. The present sequence is CC one such oligonucleotide. The oligonucleotides and the peptides encoded CC by them may be used in the prevention, diagnosis and treatment of CC diseases associated with inappropriate expression of the proteins listed CC above. Disorders that may be prevented, diagnosed and/or treated include CC multifactorial diseases with a genetic component, such as autoimmune CC diseases (e.g. rheumatoid arthritis, multiple sclerosis, diabetes, CC systemic lupus erythromatosus and Grave's disease), inflammation, cancer CC (e.g. cancers of the bladder, brain, breast, colon and kidney, landscaped of the proteins of the bladder of the nervous system and an infection of pathogenic crossing.
                                         Query Match
Best Local Similarity
                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Polymorphic nucleic acids encoding e.g. amylases, oncogenes and histones, useful for diagnosing and cancer, autoimmune diseases and infections -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1; Page 2862; 4143pp; English.
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27-DEC-2000; 2000US-0173419.
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                     Conservative
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Pred. No. 7
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                   Mismatches
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                                                                                                                                   T; 0 other;
                                           .1e+04;
                                                               DB 22;
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RESULT 11
AAH89837
                                                                                                          RESULT 12
ABK07387
                                                                                                                                                                                                                                                                                       Query Match
Best Local S
Matches 33
12-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  aberrant protein expression, including cancer, amyloidosis, diabetes, Alzheimer's disease, Down's syndrome, oedema, lupus (SLE), vasculitis, glomerulonephritis, haemolytic anaemia, thrombocytopaenia, arthritis, meningitis, muscular disorders, dementia, neurological diseases, tuberous sclerosis, male infertility, hypercalcaemia, blood pressure disorders, osteoporosis, pathogenic infections, hypercholesterolaemia, obesity or autoimmunity. The present sequence is a polymorphism-containing
                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to human nucleic acids containing single nucleotide polymorphisms (SNPs). These can be used in forensic and paternity tests, and to aid in the treatment of diseases associated with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Isolated human polynucleotides containing single nucleotide polymorphisms, useful for the treatment and diagnosis of e.g. cancer, infection and diabetes \cdot
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                                             ABK07387;
                                                                                       ABK07387
                                                                                                                                                                                                                                                                                                                                                                                                                                                  oligonucleotide fragment of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; Page 282; 475pp; English
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P-PSDB; AAM00718.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      forensic test;
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                                                                                                                                                                                                                                                                                          Local Similarity 66.0
                                                                                                                                                                                                1 tcgcggccaggctgggcgcgcgggcgggtggtgttggttacgtcag 50
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    single nucleotide polymorphism; SNP;
ic test; aberrant protein expression;
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                                                                                       standard;
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                       25 G; 9 T; 0 other;
                                                                                                                                                                                                                                                                                       Score 22.8; D
Pred. No. 7.1e
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  invention.
                                                                                                                                                                                                                                                                                                                  .1e+04;
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CC mottly pr an amberzyme (cleaving RNA with an RNR triplet), a zinzyme (cleaving RNA with a YOY motif). The CD20-targetting nucleic acid is used CC cleave RNA of CD20 in the presence of a divalent cation that is CC preferably Mg^2^+. Furthermore, it may be contacted with a cell to reduce CCD20 activity of the cell and treat a patient having a condition CC associated with the level of CD20. The treatment may further comprise the CC use of one or more therapies. In particular, the CD20 targetting CC uncleic acid may be used to treat lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular non-Hodgkin's lymphoma (NHL), bulky CC low-grade or follicular NHL, lymphocytic leukaemia, HIV (human CL), CC immunodeficiency virus) associated NHL, mantle-cell lymphoma (NCL), CC immunocytoma (IMC), small B-cell lymphocytic lymphoma, immune CC imcolocytopaenia, and inflammatory arthropathy. The NGGO-targetting CC uncleic acid is used to cleave RNA of the NGGO gene in the presence of a CC divalent cation that is preferably Mg^2-. Furthermore, the nucleic acid CC may be contacted with a cell to reduce NGGO activity of the cell and CC treat a patient having a condition associated with the level of NGGO. The CC entral nervous system (CNS) injury and cerebrovascular accident (CVA, stroke), Alzheimer's disease, dementia, multiple sclerosis (MS), CC entral cherapy-induced neuropathy, amyotrophic lateral sclerosis (MS), CC entral cherapy-induced neuropathy, amyotrophic lateral sclerosis (ALS), entral military disease, creutzfeldt-Jakob
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NGGO). The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or DNAzyme) an Inozyme (an endolytic nucleic acid cleaving a an RNA mole possessing an NCH motif), a G-cleaver (cleaving RNA with a NYN possessing an NCH motif), a G-cleaver (cleaving RNA with a NYN possessing an NCH motif), a G-cleaver (cleaving RNA with a NYN possessing an NCH motif), a G-cleaver (cleaving RNA with a NYN possessing an NCH motif), a G-cleaver (cleaving RNA with a NYN possessing an NCH motif), a G-cleaver (cleaving RNA with a NYN possessing an NCH motif), a G-cleaver (cleaving RNA with a NYN possessing an NCH motif), a G-cleaver (cleaving RNA with a NYN possessing an NCH motif), a G-cleaver (cleaving RNA with a NYN possessing an NCH motif), a G-cleaver (cleaving RNA with a NYN possessing an NCH motif), a G-cleaver (cleaving RNA with a NYN possessing an NCH motif), a G-cleaver (cleaving RNA with a NYN possessing an NCH motif), a G-cleaver (cleaving RNA with a NYN possessing an NCH motif), a G-cleaver (cleaving RNA with a NYN possessing an NCH motif), a G-cleaver (cleaving RNA with a NYN possessing an NCH motif), a G-cleaver (cleaving RNA with a NYN possessing an NCH motif), a G-cleaver (cleaving RNA with a NYN possessing an NCH motif), a G-cleaver (cleaving RNA with a NYN possessing an NCH motif), a G-cleaver (cleaving RNA with a NYN possessing an NCH motif), a G-cleaver (cleaving RNA with a NYN possessing an RNA motific RNA with a N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (RIBO-)
(BLAT/)
(MCSW/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 89;
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28-FEB-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BLATT
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; 2000US-185516P.
; 2000US-187128P.
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variants of proteins related to amylases, amyloid proteins, polymorphists related proteins, cadherin, cyclin, polymerase, oncogenes, histones, kinases, colony stimulating factors, complement related proteins, cytochromes, kinesins, cytokines, interferons, interleukins, G-protein coupled receptors and thioesterases. The present sequence is one such oligonucleotide. The oligonucleotides and the peptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of the proteins listed above. Disorders that may be prevented, diagnosed and/or treated include multifactorial diseases with a genetic component, such as autoimmune diseases (e.g. rheumatoid arthritis, multiple sclerosis, diabetes, systemic lupus erythromatosus and Grave's disease), inflammation, cancer (e.g. cancers of the bladder, brain, breast, colon and kidney,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         neuroprotective; antimicrobial; gene therapy; vaccine; amylase; cancer amyloid protein; angiopoietin; apoptosis related protein; cadherin; cyclin; polymerase; oncogene; histone; kinase; colony stimulating fact complement related protein; cytochrome; kinase; cytokine; interferon; interfeukin; G-protein coupled receptor; thioesterase; inflammation; intelfictorial disease; autoimmune disease; infection; multifactorial disease; autoimmune disease; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        disease, muscular dystrophy, and/or other neurodegenerative disease states which respond to the modulation of NOGO expression. The present sequence is a substrate sequence for a nucleic acid of the invention based on the human NOGO sequence.
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                                                                                                                                                                                                                                                  Claim 1; Page 1817; 4143pp; English.
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27-DEC-2000;
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                                                                                                                                                                                                                 present invention relates to oligonucleotides
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proteins, angiopoletin,
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                   The present invention relates to oligonucleotides encoding polymorphic CC variants of proteins related to amylases, amyloid proteins, angiopoietin, CC apoptosis related proteins, cadherin, cyclin, polymerase, oncogenes, CC histones, kinases, colony stimulating factors, complement related CC proteins, cytochromes, kinesins, cytokines, interferons, interleukins, CC G-protein coupled receptors and thioesterases. The present sequence is cone such oligonucleotide. The oligonucleotides and the peptides encoded by them may be used in the prevention, diagnosis and treatment of CC diseases associated with inappropriate expression of the proteins listed above. Disorders that may be prevented, diagnosed and/or treated include CC multifactorial diseases with a genetic component, such as autoimmune CC diseases (e.g. rheumatoid arthritis, multiple sclerosis, diabetes, systemic lupus erythromatosus and Grave's disease), inflammation, cancer (e.g. cancers of the bladder, brain, breast, colon and kidney, laukaemia), diseases of the nervous system and an infection of pathogenic
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Polymorphic nucleic acids encoding e.g. amylases, oncogenes and histones, useful for diagnosing and cancer, autoimmune diseases and infections -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAL34308 standard; DNA;
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Pred. No. 8.7e+04;
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          The present invention relates to 1,3-thiazole compounds or their salts (with position 5 substituted by 4-pyridyl group having a substituent including no aromatic group, at a position adjacent to the N of pyridyl group). The compounds of the invention are excellent inhibitors of p38 MAP kinase, phosphodiesterase IV (PD) and tumour necrosis factor (TNF) alpha production. They are antagonists of ademosine receptor. They are used for the prevent and treatment of cytckine-mediated disease, adenosine receptor and treatment of cytckine-mediated disease, autroimmune receptor they are used for the prevent and treatment of cytckine-mediated disease, adenosine receptor and treatment of cytckine-mediated disease, autroimmune the production of the prevent and treatment of cytckine-mediated disease, autroimmune the production of the prevent and treatment of cytckine-mediated disease, autroimmune the production of the prevent and treatment of cytckine-mediated disease, autroimmune the production of the prevent and treatment of cytckine-mediated disease, autroimmune the production of the prevent and treatment of cytckine-mediated disease, autroimmune the production of the prevent and treatment of cytckine-mediated disease, autroimmune the production of the prevent and treatment of cytckine-mediated disease.
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                                                                                                                                                                                                                                                            New 1.3-thiazole compounds (I) or their salts (with position substituted by 4-pyridy1) for prevention and treatment of distincluding cytokine-mediated disease, adenosine receptor-mediated diseases or inflammation .
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02-FEB-2001;
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US/08749852 NDY L. ANGUS T. EBRALD R. UMOR SUPPRESSOR 59 1. PANDERHYE P.C. GLEBE ROAD	tch al Similarity 76.9%; Pred. No. 3.8e+03; 30; Conservative 0; Mismatches 9 gccgccagcccaaccagcagcggcatgggcagcggcagc 43 [ONUCLEOTIDE NVERTING ENZ 162,484 ,661	ALIGNMENTS	PCT - US93 - 12388 - 183 PCT - US93 - 12388 - 184 US - 08 - 883 - 795A - 18 US - 08 - 883 - 795A - 18 US - 08 - 183 - 795A - 18 US - 08 - 171 - 389 - 182 US - 08 - 171 - 389 - 182 US - 08 - 475 - 228A - 182 US - 08 - 482 - 2080A - 182 US - 08 - 482 - 2080A - 182 US - 08 - 482 - 238A - 182 US - 08 - 482 - 238A - 182 US - 07 - 936 - 110 - 5 US - 07 - 936 - 110 - 5 US - 08 - 777 - 918 - 5 US - 09 - 30 - 382 - 21 US - 09 - 33 - 764 - 9 US - 08 - 531 - 927B - 17 US - 07 - 854 - 603 - 37 US - 07 - 854 - 603 - 37 US - 07 - 854 - 603 - 37 US - 08 - 171 - 389 - 376 US - 08 - 171 - 389 - 376 US - 08 - 171 - 389 - 376
	Length 42; ; Indels 0; Gaps 0	COMPOSITIONS TARGETED TO YME MRNA AND METHODS OF USE		Sequence 183, App Sequence 184, App Sequence 18, App11 Sequence 182, App Sequence 5, App11 Sequence 5, App11 Sequence 21, App1 Sequence 37, App1 Sequence 376, App Sequence 455, App

STATE:

VIRGINIA

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RESULT 3
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Best Local Similarity
Matches 28; Conserv
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TELEFAX: (703) 816-4100
INFORMATION FOR SEQ ID NO: 35:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REGISTRATION NUMBER: 32,955
REFERENCE/DOCKET NUMBER: 1579-104
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 816-4000
                                        COMPUTER: IBM PC COMPATIBLE

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/07/885,689A

FILING DATE: 19-MAY-1992

CLASSIFICATION: 435

ATTORNEY/AGENT INFORMATION:

NAME: COLORED INFORMATION:
                                                                                                                                                                           ZIP: 22040-0747
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                              STREET: P.O. Box 747
CITY: Falls Church
STATE: Virainia
                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
APPLICANT:
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LENGTH: 50 base pairs
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MEDIUM TYPE: Floppy disk
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TITLE OF INVENTION: Hormone Using a Synthetic
FITTE OF SEQUENCES: 38
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
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STRANDEDNESS: sing
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                                                                                                                                                                                                                                COUNTRY: USA
              NAME: Svensson, Leonard R. REGISTRATION NUMBER: 30,330
REFERENCE/DOCKET NUMBER:
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Park, Young
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Pred. No. 1.4e+04;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 26; Conserv
                                                                                                                                          CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/402,253
FILING DATE: 10-MAR-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/438,861
FILING DATE: 10-MAY-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: 703-241-2848 INFORMATION FOR SEQ ID NO:
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APPLICANT:
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LENGTH: 45 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: L1, Ronghao
TITLE OF INVENTION: RECEPTOR ACTIVATION BY GAS6
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genericch, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION: TELEPHONE: 703-241-1300
                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION UMBER: U5/08/628,747 FILING DATE: 17-APR-1996
                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: 1
MOLECULE TYPE:
                                              REFERENCE/DOCKET NUMBER: P9
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-225-1994
                                                                                                                             FILING DATE: 10-MAY-1995
ATTORNEY/AGENT INFORMATION:
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OTHER INFORMATION:
OTHER INFORMATION:
OTHER INFORMATION:
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               TELEPHONE: 650 -- 1-7168
                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: One DNA Way
CITY: South San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS:
                                                                                               REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
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6169070
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mark, Melanie
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SEQ ID NO:
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/note= "U7 oligonucleotide portion of synthetic
BGH gene, Figure 1."
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23:
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Pred. No. 2.1e+04;
0; Mismatches 8
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; DESCRIPTION:
US-08-628-747-23
                                                                                      US-08-766-014-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 25, Application US/08766014 Patent No. 5744312
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Best Local Similarity
Matches 28; Conserv
Query Match
Best Local Similarity 65.7
Matches 30; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               -08-766-014-25
                                                                                                                                                                                              PRIOR APPLICATION DATA: including a prior Application Data: described brior Application Number: 60/008,688 FILING DATE: December 15, 1995 ATTORNEY/AGENT INFORMATION:
NAME: WALDUTG, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 223/104
TELECOMMUNICATION INFORMATION:
TELEPAN: (213) 489-1600
TELEFAX: (213) 955-0440
TELEFAX: (213) 955-0440
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                                                                                                                                               INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 50 base pairs
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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
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LENGTH: 50 base pairs
TYPE: nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
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                                                                                                                  STRANDEDNESS: single
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California
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                                                                                                  linear
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3 West Fifth Street
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/desc = "oligonucleotide"
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                   1.18;
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   Score 20.4; DB 1; Pred. No. 3.1e+04; 0; Mismatches 16
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Pred. No. 2.6e+04;
0; Mismatches 12
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Query Match
Best Local Similarity
Vatches 25; Conserve
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                                                                                                                  RESULT 7
US-07-834-539A-50
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                                                                                  Sequence 50, Application US/07834539A GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: 202-783-6031
INFORMATION FOR SEQ ID NO:
APPLICANT: Lonberg, N
APPLICANT: Kay, Robes
TITLE OF INVENTION: I
TITLE OF INVENTION: I
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 37 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION:
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NAME: Saxe, Stephen A.
REGISTRATION NUMBER: 38,
                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE:
DESCRIPTION:
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                                                                                                                                                                                                    841 catcaccgcagatcccatggactggagccccag 873
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/0 FILING DATE: 21-APR-1995 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                   STRANDEDNESS:
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TELEFAX: 202-783-6031
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ZIP: 20004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Patino-Martin, Cristina
Kalo-Koenova, Eliza I.
del Moral-Juarez, Catalina
Faus-Santasusana, Ignacio
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                                                                Lonberg, Nils
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                                                                                                                                                                                                                                                                                                                                                                                   linear
                                                                                                                                                                                                                                                                                                                                                  other nucleic acid
/desc = "Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                 single
                                                    Robert M.
                                                                                                                                                                                                                                                                1.1%;
 Transgenic Non-Human Animals Capable of Producing Heterologous Antibodies : 77
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                                                                                                                                                                                                                                                                Score 20.2;
Pred. No. 3.
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CORRESPONDENCE ADDRESS:
ADDRESSEE: William M. Smith

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RESULT 8
US-08-053-131-67
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Best Local Similarity
Matches 25; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 67, Application US/08053131 Patent No. 5661016
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SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Lonberg, N115
APPLICANT: Kay, Robert M.
APPLICANT: Kay, Robert M.
TITLE OF INVENTION: Transgenic No. 5661016-Human Animals for
TITLE OF INVENTION: Producing Heterologous Antibodies
TITLE OF SEQUENCES: 197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/834,539A
FILING DATE: 1992-02-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M.
REGISTRATION NUMBER: 30,223
                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
CORTUMNED: PC-TOS/MS-DOS
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TELEPHONE: 415-543-9600
TELEFAX: 415-543-5043
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                                                                                                             SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: U$208/053,131
FILING DATE: 26-APR-1993
                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 197
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend Khourle and Crew
ADDRESSEE: One Market Plaza, Steuart Tower, Suite 200
                 FILING DATE: 16-DEC-PRIOR APPLICATION DATA:
                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  625 cagcagctgggcagccaaggcccctggggccag 657
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STRANDEDNESS:
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CLASSIFICATION:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                               CLASSIFICATION:
                                                                                                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                                      CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE:
APPLICATION NUMBER:
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                                     16-DEC-1992
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                                                       US 07/990,860
 US 07/810,279
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Pred. No. 3.3e+04;
0; Mismatches 8
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US-08-645-641-67
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TELEPHONE: 415-326-2400
TELEFAX: 415-326-2422
INFORMATION FOR SEQ ID NO: 67:
SEQUENCE CHARACTERISTICS:
LENGTH: 42 base pairs
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Best Local Similarity 75.8
Matches 25; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 67,
Patent No. 5
                                                                                                                                       APPLICATION NUMBER: US 07/904,068
FILING DATE: 23-JUN-1992
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M.
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 14643-0009
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-326-2400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Lonberg, Nils
APPLICANT: Kay, Robert M.
TITLE OF INVENTION: Transge
TITLE OF INVENTION: Product
NUMBER OF SEQUENCES: 150
                                                                                                         TELEFAX: 415-326-2422
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 18-MAR-1992
ATTORNEY/AGENT INFORMATION:
NAME: SMith, William M.
REGISTRATION NUMBER: 30,223
                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version
CURRENT APPLICATION DATA:
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APPLICATION NUMBER:
                                                                                    SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/645,641 FILING DATE: 20-MAY-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: USA
ZIP: 94111-3834
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STATE:
               TOPOLOGY:
                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: San Francisco
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                                STRANDEDNESS:
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                                                                     LENGTH:
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5719032
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                                nucleic acid
DEDNESS: single
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Two Embarcadero Center, 8th Floor
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DNA (primer)
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Pred. No. 3.3e+04;
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NUMBER OF SEQUENCES:

210

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; TOPOLOGY: linear; MOLECULE TYPE: DNA (primer) US-07-853-408B-67
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; Sequence 67, Application US/07853408B
; Patent No. 5789650
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                                                                                                                 US-08-096-762-67
Sequence 67 Application US/08096762
Patent No. 5814318
GENERAL INFORMATION:
APPLICANT: Lonberg, Nils
APPLICANT: Kay, Robert M.
TITLE OF INVENTION: Producing Here
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Best Local
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Best Local
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ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M.
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 1464
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-326-2400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: William M. Smith
STREET: One Market Plaza, Steuart Tower, Suite 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/853,408B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Transgenic No. 5789650-
TITLE OF INVENTION: Producing Heterologous
NUMBER OF SEQUENCES: 150
                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 42 base pairs
TYPE: NUCLEIC ACID
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CITY: San Francisco
STATE: California
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les 25; Conserv
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19920318
N: pcc
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   Transgenic No. 5814318-Human Animals for Producing Heterologous Antibodies
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75.8%;
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Pred. No. 3
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                                                                                                                                                                                                                                                                               Length 42;
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US-08-800-353-50
                                                                                                                                                                                                         Sequence 50, Application US/08800353 Patent No. 5874299
                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
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INFORMATION FOR SEQ ID NO:
                                                                      APPLICANT: Lonberg, Nils
APPLICANT: Kay, Robert M.
TITLE OF INVENTION: Transgenic |
TITLE OF INVENTION: Producing H
NUMBER OF SEQUENCES: 77
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US 07/990,860
FILING DATE: 16-DEC-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/904,068
FILING DATE: 23-JUN-1992
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 07/853,408
FILING DATE: 18-MAR-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/810,279
FILING DATE: 17-DEC-1991
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/053,131
FILING DATE: 26-APR-1993
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: 14
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-326-2400
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                   625 cagcagctgggcagccaaggcccctggggccag 657
STREET: One run.
CTTY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: FILING DATE: 22-JUI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE:
                                                                                                                                                                                                                                                                                                                        1 CAGCAGGTGCACACCCAATGCCCATGAGCCCAG 33
                                                          ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              nucleic acid
anedNess: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             E: Townsend and Townsend Khourie and Crew One Market Plaza, Steuart Tower, Suite 200
                                    E: William M. Smith
One Market Plaza, Steuart Tower, Suite 2000
                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PatentIn Release #1.0, Version #1.25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22-JUL-1993
                                                                                                                                                                                                                                                                                                                                                                                                                   1.1%;
                                                                                                                                   Transgenic No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US/08/096,762
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                                                                                                                                                                                                                                                                                                                                                                                                                   Score 20.2;
Pred. No. 3
                                                                                                               NO. 5874299-Human Animals Capable of Heterologous Antibodies
                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                   .3e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1;
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COMPUTER READABLE FORM:

COUNTRY: USA

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; TOPOLOGY: lir
; MOLECULE TYPE:
US-08-800-353-50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-308-865-67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 67, Application US/08308865 Patent No. 5877397 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 1.1%;
Best Local Similarity 75.8%;
Matches 25; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: 415-543-5043
INFORMATION FOR SEQ ID NO: 50:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/07/834
FILING DATE: 1992-02-05
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M.
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 1464
                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: William M. Smith
STREET: One Market Plaza, Steuart Tower, Suite 2000
                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Rel
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US 07/904,068
FILING DATE: 23-JUN-1992
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M.
                                                                                                              FILING DATE: 800 CLASSIFICATION: 800 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: Transgenic No. 5877397-Human Animals for TITLE OF INVENTION: Producing Heterologous Antibodies NUMBER OF SEQUENCES: 150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Lonberg, Nils APPLICANT: Kay, Robert M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: 415-543-9600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               625 cagcagctgggcagccaaggcccctggggccag 657
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                             APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                    CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER:
                                                                             FILING DATE:
                                                                                                                                                                         APPLICATION NUMBER:
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                                                                                             us/08/145,707
                                                                                                                                                                           US/08/308,865
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Pred. No. 3.3e+04;
0; Mismatches 8
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STRANDEDUES: SINGLE
TOPOLLOGY: Linear
MOLECULE TYPE: DNA (primer)
US-08-308-865-67
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US-09-042-353-228
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TELEPHONE: 415-326-2400
TELEPAX: 415-326-2422
INFORMATION: FOR SEQ ID NO: 67:
SEQUENCE CHARACTERISTICS:
LENGTH: 42 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 1.1%;
Best Local Similarity 75.8%;
Matches 25; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                             FILING DATE: 13-MAR-1998
CLASSIFICATION: 800
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/810,279
FILING DATE: 17-DEC-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/853,408
FILING DATE: 18-MAR-1992
PRIOR APPLICATION NUMBER: US 07/904,068
APPLICATION NUMBER: US 07/904,068
FILING DATE: 23-JUN-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                          FILING DATE: 16-DEC-
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                            PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 07/990,860
FILING DATE: 16-DEC-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                             FILING DATE: 22-JUL-
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Lonberg, Nils
APPLICANT: Kay, Robert M.
TITLE OF INVENTION: Trans
TITLE OF INVENTION: Produ
                                                                                                                                          PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                625 cagcagctgggcagccaaggcccctggggccag 657
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: Two Embarca
CITY: San Francisco
STATE: California
                                                                                                                                                             APPLICATION NUMBER: US 08/053,131 FILING DATE: 26-APR-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REGISTRATION NUMBER:
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                                                            APPLICATION NUMBER:
                                                                                                                      APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          94111-3834
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255458
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 E: Townsend and Townsend and Crew LLP
Two Embarcadero Center, Eighth Floor
                                      UMBER: US 08/155,301
18-NOV-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Producing Heterologous Antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Transgenic No. 6255458-Human Animals for
                                                                                                                    US 08/096,762
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US 08/161,739
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 20.2; DB 2;
Pred. No. 3.3e+04;
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PRIOR APPLICATION DATA:

PRIOR APPLICATION DATA:

APPLICATION NUMBER: FILING DATE: 10-DE

MBER: US 08/165,699 10-DEC-1993

APPLICATION NUMBER: US 01 FILING DATE: 09-MAR-1994

US 08/209,741

FILING DATE: 03-DEC-PRIOR APPLICATION DATA:

03-DEC-1993

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; STRANDEDNESS: si
; TOPOLOGY: linear
; MOLECULE TYPE: DNA
US-09-042-353-228
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REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 014643-009040US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEPAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 228:
SEQUENCE CHARACTERISTICS:
LENGTH: 42 base pairs
                                                                                                                                                                                                                                                                                                                                                                                   US-08-758-417A-76
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 밁
                                                                                                                                                                                                                                                                                                                          Sequence 76, Application US/08758417A Patent No. 6300129
GENERAL INFORMATION:
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Best Local Similarity 75.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: WO PCT/US97/21803
FILING DATE: 01-DEC-1997
ATTORNEY/AGENT INFORMATION:
NAME: APPLE, Randolnh "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: FILING DATE: 02-DEC-PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US 08/728,463 FILING DATE: 10-OCT-1996 PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: WO PCT/US96/16433 FILING DATE: 10-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US 08/544,404 FILING DATE: 10-OCT-1995
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FILING DATE: 07-DEC-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
                                                                                                                                                                                                                                 Kay, Robert M.
TITLE OF INVENTION: Transgenic No. 6300129-Human Animals
Producing Heterologous Antibodies
                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                         APPLICANT: Lonberg, Nils
                                                                                                                                 STATE: California
                                                                                              ZIP: 94111-3834
                                                                                                            COUNTRY: USA
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Pred. No. 3.3e+04;
0; Mismatches 8
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; MOLECULE TYPE: DNA
; SEQUENCE DESCRIPTION: SEQ ID NO: 76:
US-08-758-417A-76
                                                                               Query Match
Best Local S
Matches 25
                                                                                                                                                                                                                                                                                                                                       TELEFAX: (415) 576-0:
INFORMATION FOR SEQ ID NO: 76:
SEQUENCE CHARACTERISTICS:
                      625 cagcagctgggcagccaaggcccctggggccag 657
1 CAGCAGGTGCACACCCAATGCCCATGAGCCCAG
                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Serafini, Andrew T.
REGISTRATION NUMBER: 41,303
REFERENCE/DOCKET NUMBER: 014643-009030US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/728,463
FILING DATE: 10-OCT-1996
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APPLICATION NUMBER: US/08/758,417A
FILING DATE: 02-Dec-1996
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 08/096,762
FILING DATE: 22-UUI-1993
APPLICATION NUMBER: US 08/053,131
FILING DATE: 26-APR-1993
APPLICATION NUMBER: US 07/990,860
FILING DATE: 16-DEC-1992
ATTORNEY/AGENT INFORMATION:
                                                                                 ch 1.1%;
l Similarity 75.8%;
25; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 08/161,739
FILING DATE: 03-DEC-1993
APPLICATION NUMBER: US 08/155,301
FILING DATE: 18-NOV-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US 08/352,322
FILING DATE: 07-DEC-1994
APPLICATION NUMBER: US 08/209,741
FILING DATE: 09-MAR-1994
APPLICATION NUMBER: US 08/165,699
FILING DATE: 10-DEC-1993
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                                                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                       LENGTH: 42 base pairs
                                                                                                                                                                                                                                                                                                                                                                                      576-0300
                                                                                    0,
                                                                                 Score 20.2; DB 4;
Pred. No. 3.3e+04;
0; Mismatches 8;
                                                                                    8
                                                                                                                          Length 42;
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  Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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ALIGNMENTS

•	FEATURES source			TITLE JOURNAL COMMENT	REFERENCE AUTHORS	ACCESSION VERSION KEYWORDS SOURCE ORGANISM	R07222 LOCUS DEFINITION
/organisms*.nomo sapiens*. /db_xref="dD#:478966" /db_xref="taxon:9606" /clone="IMAGE:126805"	Seq primer: M13RP1 High quality sequence Location/Qual 147	Insert Size: 1185 High qality sequence stops: 1 High qality sequence stops: 1 Source: IMAGE Consortium, LLNL This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Trace considered overall poor quality Insert Length: 1185 Std Error: 0.00	Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108 Tel: 314 286 1800 Fax: 314 286 1810 Email: est@watson.wustl.edu	The WashU-Merck EST Project Unpublished (1995) Contact: Wilson R.	Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 47) 1 (bases 1 to 47) Hillier,L., Clark,N., Dubuque,T., Elliston,K., Hawkins,M., Holman,M., Kucaba,T., Le,M., Lennon,G., Marra,M., Parsons,J., Parsickar T. Polificar B. Corpo.	R07222.1 GI:759145 EST. human. Homo sapiens	

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Washington University School of MedicineP
Washington University School of Medic
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Marra, M., Hillter, L., Allen, M., Bowles, M., Dietrich, N., Dubuque, T. Geisel, S., Kucaba, T., Lacy, M., Le, M., Martin, J., Morris, M., Schellenberg, K., Steptoe, M., Tan, F., Underwood, K., Moore, B., Theising, B., Wylle, T., Lennon, G., Soares, B., Wilson, R. and Waterston, R.
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While, T., Lennon, G., Soares, B., Wilson, R. and Waterston, R.
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40 bp mRNA linear EST 06-FEB-1998 vt78d02.rl Barstead mouse irradiated colon MPLRB7 Mus musculus cDNA clone IMAGE:1177251 5' similar to SW:ETS4_DROME P29775 DNA-BINDING
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: mouseest@watson.wustl.edu
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
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/notc-"Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site_1: EcoRI; Site_2: NotI; Tissue obtained from 8 week old mouse. Colon was harvested 72 hours after the colon was barvested 72
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/lab_host="DH10B (ampicillin resistant)"
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/sex="male"
                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Mus musculus"
/strain="FVB/N"
/db_xref="taxon:10090"
/clone="IMAGE:1177251"
                                                                                                                                                                                                                                                                         /clone_lib="Barstead mouse irradiated colon MPLRB7"
/dev_stage="8 weeks"
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RESULT AA789732

DEFINITION

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AU104284
AU104284 Sugano Homo sap
HEP07220, mRNA sequence.
AU104284
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
Email: ysuzukleims.u-tokyo.ac.jp
Suzuki.y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sugano,S. Construction and characterization of a full length-enriched and a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Suzuki, Y., Taira, H., Tsunoda, T., Mizushima-Sugano, J., Sese, J., Hat, Ota, T., Isogai, T., Tanaka, T., Morishita, S., Okubo, K., Sakaki, Y., Nakamura, Y., Suyama, A. and Sugano, S.
Diverse transcriptional initiation revealed by fine, large-scale mapping of mRNA start sites
EMBO Rep. 2 (5), 388-393 (2001)
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AU104281 Sugano Homo sapiens cDNA library Homo sapiens cDNA clone HEP04739, mRNA sequence.
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1 (bases 1 to 50)
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Similarity 79.5%;
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10 c 11 g 9 t
                                       Sugano Homo sapiens cDNA library Homo sapiens cDNA clone
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                                                                                                                                                                                                                                                                                                                                                                                          /organism="Homo sapiens"
/db_xref="taxon:9606"
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                                                                                                                                                                                                                                            Score 24.8;
Pred. No. 3.
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                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                        DB 9;
1.8e+05;
les 7;
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Matches 29
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                                                                     Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedal, Minatoku, Tokyo 108-8639, Japan
Email: ysuzuki@ims.u-tokyo.acjp
Suzuki,Y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sugano,S. Construction and characterization of a full length-enriched and a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997).
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AU104283 Sugano Homo sapiens
HEP06519, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
Email: ysuzuki@ims.u-tokyo.ac.jp
Suzuki,Y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sugano,S. Construction and characterization of a full length-enriched and a 5'-end-enriched CDNA library. Gene 200 (1-2), 149-156 (1997).
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Diverse transcriptional initiation revealed by fine, large-scale mapping of mRNA start sites
EMBO Rep. 2 (5), 388-393 (2001)
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21270072
                                                                                                                                                                                                                                                                                  ,Y., Nakamura,Y., Suyama,A. and Sugano,S.
Diverse transcriptional initiation revealed by fine, large-scale mapping of mRNA start sites
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,H., Ota,T., Isogai,T., Tanaka,T., Morishita,S., Okub
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AU104283.1 GI:13553804
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EST.
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                                                                                                                                                                                                                                                                                                                                                                                         (bases 1 to 50)
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/db_xref="taxon:9606"
/clone="HEP07220"
/clone_lib="Sugano Homo sapiens
/lic 28 g 4 t
/organism="Homo sapiens"
/db_xref="taxon:9606"
                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                            Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                              Metazoa;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Chordata; Craniata; Vertebrata; Euteleostomi; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
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Pred. No. 3.8e+05;
0; Mismatches 7;
                                                                                                                                                                                                                                                                   (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       bp mRNA linear EST 30-AUG-200 cDNA library Homo sapiens cDNA clone
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AA108149/c
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Matches 31;
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JOURNAL
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Best Local
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1426 gcctggcccagggcctgaaacccgccctcaggggcctctctcctg 1470
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Marra M/Mouse EST Project Washington University School of MedicineP 4444 Forest Parkway, Box 8501, St. Lc Tel: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 49)
1 (bases 1 to 49)
Marra,M., Hillier,L., Allen,M., Bowles,M., Dietrich,N., Dubuque,T., Geisel,S., Kucaba,T., Lacy,M., Le,M., Martin,J., Morris,M., Schellenberg,K., Steptoe,M., Tan,F., Underwood,K., Moore,B., Theising,B., Wylie,T., Lennon,G., Soares,B., Wilson,R. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: mouseest@watson.wustl.edu
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AA108149 49 bp mRNA linear EST 03-FEB-1997 m158h07.rl Stratagene mouse testis (#937308) Mus musculus clNA clone IMAGE:516253 5' similar to gb:U29152 Mus musculus aldose reductase mRNA, complete cds (MOUSE);, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unpublished (1996)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Waterston, R.
The WashU-HHMI Mouse EST Project
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AA108149
AA108149.1 GI:1659201
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Similarity 77.8%;
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                                                                                                      Conservative
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                                                                                                                                                                                                                                                                                 /tlssue_type="testis"
/tlssue_type="testis"
/dev_stage="10-12 week old"
/lab_host="SOLR (kanamycin resistant)"
/lab_host="SOLR (kanamycin resistant)"
/note="Organ: testis; Vector: pBluescript SK-; Site_1:
/note="Organ: testis; Vector: pBluescript SK-; Site_1:
/note="Organ: testis; Vector: pBluescript Sk-; Site_1: About Site_2: NhOI; Cloned unidirectionally. Primer:
/note="Organ: testis; Vector: pBluescript Site_2: NhOI; Cloned unidirectionally. Primer:
/note="Organ: testis; Vector: pBluescript Sk-; Site_1: NhOI; Vector: pBluescript Sk-; Site_1:
/note="Organ: testis; Vector: pBluescript Sk-; Site_1: NhOI; 
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/clone_lib="Sugano
12 c 26 g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /db_xref="taxon:10090"
/clone="IMAGE:516253"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /organism="Mus musculus"
/strain="Inbred CD-1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /clone_lib="Stratagene mouse testis (#937308)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
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                                                                                                                                   1.2%;
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                                                                                                   Score 22.6; DB 9;
Pred. No. 9.6e+05;
0; Mismatches 14;
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Pred. No. 7.5e+05;
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cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequ
Clone distribution: NCI-CGAP clone distribution info
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/inage/inage.hunl
Insert Length: 1032 Std Error: 0.00
Seq primer: -40UP from Gibco
High quality sequence stop: 1.
Cocation/Qualifiers
Homo sapiens
Eukaryota; Metazoa; (
Mammalia; Eutheria; 1
1 (bases 1 to 50)
                                                                                                                      AU104286 50 AU104286 Sugano Homo sapiens HEP15925, mRNA sequence.
AU104286 AU104286.1 GI:13553807
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46 bp mRNA linear EST 14-DEC-19
tq12e07.x1 NCI_CGAP_Ut3 Homo sapiens cDNA clone IMAGE:2208612 3'
similar to SW:PRPL_HUMAN P10162 SALIVARY PROLINE-RICH PROTEIN PO
                                                                                                         EST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: cgapbs-r@mail.nlh.gov
Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Contact: Robert Strausberg, Ph.D.
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 1.45 kb. Life Technologies catalog #:
11541-018"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /Clone="IMAGE:2208612"
/Clone_lib="NOI_CGAP_Ut3"
/tissue_type="poorly-differentiated endometrial adenocarcinoma, 2 pooled tumors"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="Homo sapiens"
/db_xref="taxon:9606"
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                   Chordata;
Primates;
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Diverse transcriptional initiation revealed by fine, large-scale mapping of mRNA start sites
EMBO Rep. 2 (5), 388-393 (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: ysuzuki@ims.u-tokyo.ac.jp
Suzuki,Y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sugano
,S. Construction and characterization of a full length-enriched and
a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997).
                                                                                                                                         Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0171 row: A column: 16
Seq primer: CACACAGGAACAGCTATGACC
Class: plasmid ends
                                                                                                                                                                                                                                                                                                              University of Utah (University of Utah Rm. 308, Biomedical
                                                                                                                                                                                                                                                                                                                                                                                                                                  Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D., Weiss,R. Mouse whole genome scaffolding with paired end reads from 10kb mouse whole genome scaffolding with paired end reads from 10kb
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1M0171A16R Mouse 10kb plasmid UUGC1M
clone UUGC1M0171A16 R, DNA sequence.
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Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
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                                                                                                                                                                                                                                                 Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                               Unpublished (2000)
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/db_xref="taxon:9606"
/clone="HEP15925"
/organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0171A16"
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Pred. No. 1.5e+06;
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                                                                                                                                                                                                                                Contact: Yutaka Suzuki
Department of Virology
Institute of Medical Science, University of Tokyo
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
Email: ysuzuki@ims.u-tokyo.ac.jp
Suzuki,Y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sugano,S. Construction and characterization of a full length-enriched and a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997).
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Diverse transcriptional initiation revealed by fine, large-scale mapping of mRNA start sites
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HEP05158, mRNA sequence.
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1 (bases 1 to 50)
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                                                                                   /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="HEP05158"
                                 /clone_lib="Sugano Homo sapiens cDNA library'
11 c 18 g 10 t
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Pred. No. 1.6e+06;
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CDNA library Homo sapiens cDNA clone
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ACCESSION VERSION

SOURCE KEYWORDS

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REFERENCE

AUTHORS

DEFINITION AU104319

Snoo RESULT

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MEDLINE JOURNAL TITLE

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BASE COUNT ORIGIN

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                                                       AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedal, Minatoku, Tokyo 108-8639, Japan
Email: ysuzuki@ims.u-tokyo.ac.jp
Suzuki,Y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sugano,S. Construction and characterization of a full length-enriched and a 5'-end-enriched CONA Library. Gene 200 (1-2), 149-156 (1997).
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Diverse transcriptional initiation revealed by fine, large-scale mapping of mRNA start sites

EMBO Rep. 2 (5), 388-393 (2001)
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AU107649 Sugano Homo sapiens cDNA library Homo sapiens cDNA clone
HSIO2001, mRNA sequence.
AU107649
1 (bases I to 50)
Suzuki,Y., Taira,H., Tsunoda,T., Mizushima-Sugano,J.,
Ota,T., Isogai,T., Tanaka,T., Morishita,S., Okubo,Y., Nakamura,Y., Suyama,A. and Sugano,S.
Diverse transcriptional initiation revealed by fine, 1
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AU107650.1 GI:13557171
EST.
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AU107650 Sugano Homo sapiens
HSI03379, mRNA sequence.
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Mammalia; Eutheria; Primates;
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Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/db_xref="taxon:9606"
/clone="HST02001"
/clone_lib="Sugano Homo s:
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                                                                                                                                                                                                                                                                                   Contact: Yutaka Suzuki
Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
Email: ysuzuki@ims.u-tokyo.ac.jp
Suzuki,Y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sugano,S. Construction and characterization of a full length-enriched and a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997).
Location/Qualifiers
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Institute of Medical Science, University of Tokyo
1. Stitute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
Email: ysuzuki@ims.u-tokyo.ac.jp
Suzuki,Y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sugano
Suzuki,Y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sugano
5. Construction and characterization of a full length-enriched and
a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997).
Location/Qualifiers
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EMBO Rep. 2 (5), 388-393 (2001)
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EMBO Rep. 2 (5), 388-393 (2001)
21270072
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1 (Dases 1 to 50)

1 (Dases 1 to 50)

Suzuki,Y., Taira,H., Tsunoda,T., Mizushima-Sugano,J., Sese,J., Haira,T., Taona,T., Morishita,S., Okubo,K., Sakaki,Y., Nakamura,Y., Suyama,A. and Sugano,S.

Diverse transcriptional initiation revealed by fine, large-scale
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AU107651 Sugano Homo sapiens
HSI07783, mRNA sequence.
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/db_xref="taxon:9606"
/db_xref="HSI03379"
/clone="HSI03379"
/clone_1lb="Sugano Homo sapiens cDNA library"
/clone_1lb="Sugano Homo sapiens cDNA library"
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/db_xref="taxon:9606"
/clone="HS107783"
/clone=11b="Sugano Homo sapiens cDNA library"
/clone_11b="Sugano Homo sapiens cDNA library"
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Pred. No. 1.9e+06;
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Pred. No. 1.9e+06;
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CASO1643, mRNA sequence.
AU104063
AU104063.1 GI:13553584
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Department of Virology
Institute of Medical Science, University of Tokyo
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
Email: ysuzuki@ims.u-tokyo.ac.jp
Suzuki@y. Yoshitomo-Nakagawa.K., Maruyama,K., Suyama,A. and Sugano
,S. Construction and characterization of a full length-enriched and
a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997).
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Diverse transcriptional initiation revealed by fine, large-scale mapping of mRNA start sites
                                    Suzuki,Y., Taira,H., Tsunoda,T., Mizushima-Sugano,J., Sese,J.,
H., Ota,T., Isogai,T., Tanaka,T., Morishita,S., Okubo,K., Saka
,Y., Nakamura,Y., Suyama,A. and Sugano,S.
Diverse transcriptional initiation revealed by fine, large-scal
mapping of mRNA start sites
EMBO Rep. 2 (5), 388-393 (2001)
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AU104502 Sugano Homo sapiens
ERV6C934, mRNA sequence.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
Contact: Yutaka Suzuki
Department of Virology
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Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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78.18;
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Pred. No. 2.1e+06;
0; Mismatches 7;
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Search completed: August 6, 2002, 21:35:03 Job time: 6147 sec
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                                                                                                                          501 agaaggcggcagcgggggcagtgggtctcgagaga 535
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4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
Email: ysuzuki@ims.u-tokyo.ac.jp
Suzuki,Y., yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sugano,S. Construction and characterization of a full length-enriched and a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997).
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Post-processing: Minimum Match 0%
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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19	٠		41	35	US-09-953-198-123	123, A
			44	æ	US-08-472-801-999	999, 1
c 21			44	10	US-08-668-235-1000	a
22	٠		48	37	US-10-017-974-22699	
23	22	1,2	22	14	US-09-064-645-23	
24	22	•	22	22	2-5	Sequence 23, App.
25	22		45	<u>υ</u>	US-09-959-356A-7	Sequence 7, Appli
c 26	22		45	<u>ω</u>	US-09-959-356A-8	
27	22		48	ω	•	Sequence 3326, Ap
28	22		48	37	017-	Sequence 34757,
	22		50	29	-09-755	Sequence 17016, A
c 30	21.8		49	64	-60-253-	9804, A
	<u>2</u>	,	;		-00-755	1

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US-09-570-593-11/c ; Sequence 11, Application US/09570593 ; GENERAL INFORMATION:
                    ; PRIOR APPLICATION NUMBER: 60/13; PRIOR FILING DATE: 1999-05-14; NUMBER OF SEQ ID NOS: 13; SOFTWARE: FASTSEQ for Windows V SEQ ID NO 10; LENGTH: 31
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                                                                                                                                                                                                                            US-09-570-593-10
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SEQ ID NO 11
LENGTH: 34
                                                                                                                                                                                                     Sequence 10, Application US/09570593 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                  Query Match 1.8%;
Best Local Similarity 100.0%;
Matches 34; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: 2300-1556
CURRENT APPLICATION NUMBER: US/09/570,593
CURRENT FILING DATE: 2000-05-12
PRIOR APPLICATION NUMBER: 60/134,112
PRIOR FILING DATE: 1999-55-14
NUMBER OF SEQ ID NOS: 13
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                          APPLICANT: Kaufmann, Joerg
APPLICANT: Xin, Hong
APPLICANT: Xin, Hong
APPLICANT: Xin, Hong
APPLICANT: Xin, Hong
APPLICANT: MXIn, Hong
APPLICANT: Harrowe, Greg
TITLE OF INVENTION: EXPRESSION OF ETS-DOMAIN PROTEINS IN
TITLE OF INVENTION: CANCER
FILE REFERENCE: 2300-1556
CURRENT APPLICATION NUMBER: US/09/570,593
CURRENT FILING DATE: 2000-05-12
CURRENT FILING DATE: 2000-05-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: HATTOWE, Greg TITLE OF INVENTION: EXPRESSION OF ETS-DOMAIN PROTEINS IN TITLE OF INVENTION: CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
OTHER INFORMATION: GSEF-specific primer
ORGANISM: PArtificial Sequence
                                                                                                                                                                                                                                                                                                 1850 ccaggagtgcccccgggaatggataataaagata 1883
                                                                                                                                                                                                                                                                                  34 CCAGGAGTGCCCCCGGGAATGGATAATAAAGATA 1
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21.6
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                                               FastSEQ for Windows Version 4.0
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448
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US-09-216-745-4273
US-09-30-423-3875
US-09-755-374A-23624
US-09-755-374A-23626
US-09-848-754A-7792
US-09-848-754A-7792
US-09-848-754A-791
US-09-214-913-51
US-09-718-159-3
US-09-718-159-3
US-09-718-159-3
US-09-718-159-3
US-09-533-5808-45
US-09-771-372-8
                                                                                                                                                                                                                                                                                                                                   0;
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Pred. No. 1.4e+0
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                                                                                                                                                                                                                                                                                                                                    Mismatches
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Sequence
Sequence
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3875, Ap
23624, A
23624, A
23626, Ap
7792, Ap
3298, Ap
3298, Ap
3298, Ap
11, Appl
11, Appl
11, Appl
11, Appl
11, Appl
145, Appl
15, Appl
16, Appl
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; LOCATION: (26)...(0)
OTHER INFORMATION: 2 of 2 allelic variants (3339 is other entry);
NAME/KEY: misc_feature
LOCATION: (25)...(26)
OTHER INFORMATION: Nucleotide deleted between bases 25 and 26
NAME/KEY: misc_feature
LOCATION: (0)...(0)
COTHER INFORMATION: Accession number cg44930423
US-09-755-374A-3340
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; OTHER INFORMATION: GSEF-specific primer US-09-570-593-10
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                                                    CURRENT APPLICATION NUMBER: US/09/755,374A
CURRENT FILING DATE: 2001-01-08
PRIOR APPLICATION NUMBER: 60/174962
PRIOR FILING DATE: 2000-01-07
NUMBER OF SEQ ID NOS: 28742
SEQ ID NO 26208
LENGTH: 50
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LENGTH: 50
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Best Local Similarity
Matches 31; Conserv
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Best Local
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CURRENT ETLING DATE: 2001-01-08
PRIOR APPLICATION NUMBER: 60/174962
PRIOR FILING DATE: 2000-01-07
NUMBER OF SEQ ID NOS: 28742
                                                                                                                                                                                   APPLICANT: Leach, Martin
APPLICANT: Shinkets, Richard A.
APPLICANT: Shinkets, Richard A.
TITLE OF INVENTION: Nucleic Acids Containing
TITLE OF INVENTION: Use Thereof
FILE REFERENCE: 15966-611
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Nucleic Acids Containing Single Nucleotide Polymorphisms TITLE OF INVENTION: Use Thereof FILE REFERENCE: 15966-611
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Leach, APPLICANT: Shimk
                                    TYPE: DNA
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ORGANISM: Homo sapiens
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                ORGANISM: Homo sapiens
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FEATURE:
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Conservative 0;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 24.4; DB 29;
Pred. No. 9.1e+05;
0; Mismatches 16;
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5e+04;
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; NAME/KEY: misc_feature
; LOCATION: (25)...(26)
; OTHER INFORMATION: Nucleotide de
; NAME/KEY: misc_feature
; LOCATION: (0)...(0)
; OTHER INFORMATION: Accession nun
US-09-755-374A-2825
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LOCATION: (25)...(26)
OTHER INFORMATION: Nucleotide deleted between
NAME/KEY: misc_feature
LOCATION: (0)...(0)
OTHER INFORMATION: Accession number cg44009035
US-09-755-374A-26208
                                                                                                                                             RESULT 6
US-10-017-974-35747
; Sequence 35747, Application US/10017974
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Shimkets, Richard A.
TITLE OF INVENTION: Nucleic Acids Containing Si:
TITLE OF INVENTION: Use Thereof
FILE REFERENCE: 15965-611
CURRENT APPLICATION NUMBER: US/09/755,374A
CURRENT TLING DATE: 2001-01-08
PRIOR FILING DATE: 2000-01-07
PRIOR FILING DATE: 2000-01-07
NUMBER OF SEQ ID NOS: 28742
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LENGTH: 50
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Best Local Similarity 71.1
Matches 32; Conservative
                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Blatt, Lawrence
APPLICANT: McSwiggen, James
TITLE OF INVENTION: Nucleic Acid-Based Treatment of Diseases
TITLE OF INVENTION: Nile Virus Infection
FILE REFERENCE: MBHB00,1109-A (400/037)
CURRENT APPLICATION NUMBER: US/10/017,974
CURRENT FILING DATE: 2001-12-10
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OTHER INFORMATION: 1 of 2 allelic
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LOCATION: (26)...(0)
OTHER INFORMATION: 2 of 2
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                                                                                                                                                                                                                                                                                                                           Conservative
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71.1%;
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Pred. No. 9.9e+05;
0; Mismatches 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                  number cg43969816
                                                                                                                                                                                                                                                                                                                         Score 23.8; DB 29;
Pred. No. 1.2e+06;
0; Mismatches 12;
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                                                                        or Conditions Related
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; OTHER INFORMATION: Description of Artificial Sequence: Enzymatic Nucleic Acid US-10-017-974-35747
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US-09-422-978-175
                                                                                                                                                                                       US-09-718-159-5; Sequence 5, Applicat; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; LOCATION: 24
; OTHER INFORMATION: 99-13320-352 : polymorphic base
US-09-422-978-175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/422,978
CURRENT FILING DATE: 1999-10-20
EARLIER APPLICATION NUMBER: US 09/298,850
EARLIER FILING DATE: 1999-04-21
EARLIER APPLICATION NUMBER: US 60/109,732
EARLIER APPLICATION NUMBER: US 60/082,614
EARLIER APPLICATION NUMBER: US 60/082,614
EARLIER FILING DATE: 1998-04-21
EARLIER FILING DATE: 1998-04-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 37080
SOFTWARE: PatentIn version 3
SEQ ID NO 35747
LENGTH: 48
TYPE: RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 11796
SEQ ID NO 175
                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local :
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Best Local Similarity
Matches 29: Conserv
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APPLICANT: Blumenfeld, Marta
APPLICANT: Chumakov, Ilya
TITLE OF INVENTION: Biallelic markers for
FILE REFERENCE: GENSET.020CP1
                                                                                                                                                                        APPLICANT:
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                                                                  APPLICANT:
                                                                                     APPLICANT:
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                  TITLE OF INVENTION:
                                     APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Homo Sapiens FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 47
                                                                                                                                                                                                                                                                                                             1484 agccaggccctgagatgggggaaaaacgggcagtgtgctctgctgct 1529
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                                                                                                                                                                                                                                                                                           1 aggcaagccctgtaccctcgggargccggctgtgtgctctgctgtt 46
            Parkes, Deborah
Parry, Gordon
Steinbrecher, Renate
Van Heuit, Pamela Toy
Xuan, Jian-ai
XUENTION: DNA Encoding a N
                                                                                                                                                                                                        Application US/09718159
                                                                                                                                                    Harkins, Richard
Lin, Richard
                                                                                                                  Monteclaro, Felipe
                                                                                                                                         Luke, May
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51790AUSM1
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Pred. No. 1.3e+06;
3; Mismatches 14;
                                                                                                                                                                                                                                                                                                                                                               Score 23.2; DB 18
Pred. No. 1.5e+06;
1; Mismatches 14
                  Novel
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              ETS Polypeptide
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SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 5
LENGTH: 23
TYPE: DNA
ORGANIEM: Artificial Sequence
FEATURE: INFORMATION: Description of Artificial Sequence: probe
US-09-718-159-5
                                                                                        US-09-718-159-10 ; Sequence 10, Application US/09718159 ; GENERAL INFORMATION:
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US-09-718-159-9
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Best Local Similarity
Traches 17; Conserv
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SEQ ID NO 9
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Best Local Similarity
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APPLICANT:
                              APPLICANT:
APPLICANT:
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OTHER INFORMATION: Description of Artificial Sequence: antisense-09-718-159-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Van Heuit, Pamela Toy
APPLICANT: Xuan, Jian-ai
TITLE OF INVENTION: DNA Encoding a Novel ETS Polypeptide
FILE REFERENCE: 51790AUSM1
CURRENT APPLICATION NUMBER: US/09/718,159
CURRENT FILING DATE: 2000-11-21
PRIOR APPLICATION NUMBER: 60/168,182
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APPLICANT:
APPLICANT:
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PRIOR FILING DATE: 1999-11-30
NUMBER OF SEQ ID NOS: 11
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CURRENT FILING DATE: 2000-11-21
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PRIOR FILING DATE: 1999-
NUMBER OF SEQ ID NOS: 11
 APPLICANT:
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TYPE: RNA
ORGANISM: Artificial Sequence
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Monteclaro, Felipe
Parkes, Deborah
Parry, Gordon
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Parry, Gordon
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                                             Luke, May
                                                            Harkins, Richard
Lin, Richard
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73.9%;
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100.0%; Pred. No.
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Pred. No. 1.6e+06;
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1.6e+06;
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PRIOR APPLICATION NUMBER: 60/16;
PRIOR FILING DATE: 199-11-30
NUMBER OF SEQ ID NOS: 11
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 10
LENGTH: 23
TYPE: RNA
ORGANISM: Artificial Sequence
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; OTHER INFORMATION:
US-09-718-159-10
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US-09-755-374A-9716
                  RESULT 12
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                                                                                                                                                                                                           ; OTHER INFORMATION: Description of Artificial Sequence: antisense US-09-718-159-11
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Best Local S
Matches 19
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SEQ ID NO 11
LENGTH: 23
TYPE: RNA
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Best Local Similarity
Matches 21; Conserv
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                                                                                                                                                                                                                                                                                                                                 APPLICANT: Xuan, Jian-ai
TITLE OF INVENTION: DNA Encoding a Novel ETS Polypeptide
FILE REFERENCE: 51790AUSM1
CURRENT APPLICATION NUMBER: US/09/718,159
CURRENT FILING DATE: 2000-11-21
PRIOR APPLICATION NUMBER: 60/168,182
PRIOR FILING DATE: 1999-11-30
NUMBER OF SEQ. ID NOS: 11
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CURRENT APPLICATION NUMBER: US/09/718,159
CURRENT FILING DATE: 2000-11-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Xuan, J. TITLE OF INVENTION: FILE REFERENCE: 51
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                                                                             1699 ccagcatttccagagcagagcct 1721
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                                                                1 ccagcauuuccagagcagagccu
                                                                                                                                 19;
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Parry, Gordon
Steinbrecher, Renate
Van Heuit, Pamela Toy
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Van Heuit, Pamela Toy
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                                                                                                                                               1.2%;
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                                                                                                                                             Score 23;
Pred. No.
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Pred. No. 1.6e+06;
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                                                                                                                                                            Length 23;
                                                                                                                                Indels
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Sequence 9716, Application US/09755374A GENERAL INFORMATION:
APPLICANT: Leach, Martin

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: LOCATION: (0)...(0)
: OTHER INFORMATION: Accession number cg43952512
US-09-755-374A-9716
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: GLEDN, Matthew
APPLICANT: Grigor, Murray R
TITLE OF INVENTION: Compositions isolated
TITLE OF INVENTION: tissues and methods f
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Matches
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SOFTWARE: FastSEQ for
SEQ ID NO 21580
                                                                                         Sequence 28060, Application US/09755374A GENERAL INFORMATION:
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Best Local Similarity
Matches 29; Conserv
                                                                                                                                                                                                                                                                                                            Query Match
APPLICANT: Leach, Martin
APPLICANT: Shimkets, Richard A.
TITLE OF INVENTION: Nucleic Acids
TITLE OF INVENTION: Use Thereof
FILE REFERENCE: 15966-611
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/60/253,652
CURRENT FILING DATE: 2000-11-28
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TITLE OF INVENTION: Nucleic Acids Containing
TITLE OF INVENTION: Use Thereof
FILE REFERENCE: 15966-611
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILE REFERENCE: 1055P2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: misc_feature LOCATION: (25)...(26) OTHER INFORMATION: Nucleotide
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LOCATION: (26)...(0)
OTHER INFORMATION: 2 of 2 allelic variants (9715 is other entry)
                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
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                                                                                                                                                                                                                     383 acagacacagccggccagcccaaaccagcagcggcatgggc 421
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Pred. No. 1.7e
0; Mismatches
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                                      Containing Single Nucleotide Polymorphisms and Meti
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1.7e+06
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Search completed: August Job time: 9294 sec

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2002, 23:36:06

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APPLICANT: Hesed 2
APPLICANT: Smith, Larry J.
TITLE OF INVENTION: Method and Compositions for TITLE OF INVENTION: Reprogramming
FILE REFERENCE: Hesed 2
CURRENT APPLICATION NUMBER: US/08/472,801
CURRENT FILING DATE: 1995-06-07
NUMBER OF SEQ ID NOS: 3601
SOPTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 161
LENGTH: 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; NAME/KEY: misc_feature; LOCATION: (0)...(0)
; OTHER INFORMATION: Accession number cg44009326
US-09-755-374A-28060
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                                                                                                                                                  ; ORGANISM: Homo sapiens US-08-472-801-161
                                                                                                                                                                                                                                                                                                                                                                                      US-08-472-801-161/c
: Sequence 161, Applic
: GENERAL INFORMATION:
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LENGTH: 50
TYPE: DNA
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Best Local Similarity 66.0
Matches 33; Conservative
                                                             Query Match 1.3
Best Local Similarity 75.7
Matches 28; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/755,374A
CURRENT FILING DATE: 2001-01-08
PRIOR APPLICATION NUMBER: 60/174962
PRIOR FILING DATE: 2000-01-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 28742
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: misc_feature
LOCATION: (25)...(26)
OTHER INFORMATION: Nucleotide deleted between bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LOCATION: (26)...(0)
OTHER INFORMATION: 2 of 2
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GGAGGTGCGCCCAAGCTGGAGGAGCAGGCCCAGCAG 1
                                                                                                                                                                                                                                                                                                                                                                                                        Application US/08472801
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                                                                             1.2%;
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Pred. No. 1.8e+06;
0; Mismatches 17;
                                                                             Score 22.6; DB 8;
Pred. No. 1.9e+06;
                                                                 Mismatches
                                                                                              DB 8;
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                                                                                              Length
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                                                                 0;
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                                                                 Gaps
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Minimum
Maximum
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No.
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
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21
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   Pending_Patents_NA_New:*

1: /cgn2_6/ptodata/1/pna/JCT_NEW_COMB.seq:*

2: /cgn2_6/ptodata/1/pna/JUS06_NEW_COMB.seq:*

3: /cgn2_6/ptodata/1/pna/JUS07_NEW_COMB.seq:*

4: /cgn2_6/ptodata/1/pna/JUS09_NEW_COMB.seq:*

5: /cgn2_6/ptodata/1/pna/JUS09_NEW_COMB.seq:*

6: /cgn2_6/ptodata/1/pna/JUS10_NEW_COMB.seq:*

7: /cgn2_6/ptodata/1/pna/JUS10_NEW_COMB.seq:*

8: /cgn2_6/ptodata/1/pna/JUS10_NEW_COMB.seq:*
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1894
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  GenCore version 4.5
Copyright (c) 1993 - 2000 Comp
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   В
US-60-384-980-216
US-09-745-237A-3875
US-09-90-217-51
US-09-910-212-63
US-10-126-022-268
US-10-126-306-7880
US-10-126-306-7880
US-09-718-321A-1048
US-09-745-237A-3833
US-60-384-980-69
US-10-09-926-1257
US-09-745-237A-3877
US-09-745-237A-3877
US-09-745-237A-3877
US-09-745-237A-3877
US-09-745-237A-3877
US-09-745-237A-3877
US-09-745-237A-3650
US-10-156-306-3945
US-09-718-321A-1047
US-09-718-321A-1047
US-09-718-321A-1047
US-09-718-321A-228
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        Sequence 216, App
Sequence 3875, Ap
Sequence 268, App
Sequence 268, App
Sequence 258, Appl
Sequence 25, Appl
Sequence 25, Appl
Sequence 3833, Ap
Sequence 3877, Ap
Sequence 3877, Ap
Sequence 4024, Appl
Sequence 4027, Appl
Sequence 4027, Appl
Sequence 4027, Appl
Sequence 4027, Appl
Sequence 7660, Ap
Sequence 1047, Appl
Sequence 7660, Ap
Sequence 3870, Appl
                                                                                                                                                                                                                                                                                                        Description
sequence
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US-09-745-237A-3875 US-09-745-237A-3875 US-09-745-237A-3875 Sequence 3875, Application U GENERAL INFORMATION: APPLICANT: Ribozyme Pharmac APPLICANT: Blatt, Larry APPLICANT: McSwiggen, Jim FITLE OF INVENTION: Method FILE REFERENCE: 400/007 (MB CURRENT APPLICATION NUMBER: CURRENT FILING DATE: 2007- NUMBER OF SEQ ID NOS: 4550 SOFTWARE: Patentin version SEQ ID NO 3875 LENGTH: ARA ORGANISM: Artificial Seque FEATURE: OTHER INFORMATION: Descrip US-09-745-237A-3875	Query Match Best Local Similarity Matches 29; Conserva Qy 1233 acaaggagaaaggga Db 41 AAAGGGGGAAGAGAA	RESULT 1 US-60-384-980-216/c ; Sequence 216, Application U ; GENERAL INFORMATION: APPLICANT: BRIStol-Myers S ; TITLE OF INVENTION: HUMAN FILE REFERENCE: DO211 PSP ; CURRENT APPLICATION AUMBER ; CURRENT FILING DATE: 2002 ; NUMBER OF SEQ ID NOS: 1219 ; SOFTWARE: Patentin version ; SEQ ID NO 216 ; LENGTH: 41 ; TYPE: DNA ; ORGANISM: Homo Sapiens US-60-384-980-216		26 19.4 1.0 27 19.4 1.0 28 19.4 1.0 29 19.4 1.0 29 19.4 1.0 29 19.4 1.0 29 19.2 1.0 29.3 19.2 1.0 29.3 19.2 1.0 29.3 19.2 1.0 29.3 19.2 1.0 29.3 19.2 1.0 29.3 19.2 1.0 29.3 19.2 1.0 29.3 19.2 1.0 29.3 19.2 1.0 29.3 19.2 1.0 29.3 19.2 1.0 29.3 19.2 1.0 29.3 19.3 19.3 19.3 29.3 29.3 29.3 29.3 29.3 29.3 29.3 2
S/09745237A euticals, Inc. and Reagent for the Trea HB00-918-A) US/09/745,237A 04-15 3.0 nce nce	1.2%; Score 21.8; DB 8; 70.7%; Pred. No. 3.1e+05; tive 0; Mismatches 12; tcttcaaaattgaggactcagcccag 12	CION: ION: ION: ITON: HYGER Squibb Company ION: HYMAN SINGLE NUCLEOTIDE POLYMORPHISMS D0211 PSP UTION NUMBER: US/60/384,980 DATE: 2002-06-03 D NOS: 1219 D NOS: 1219 DLIN version 3.1	ALIGNMENTS	40 7 US-10-068-965-5 41 1 PCT-US02-12063-402 41 7 US-09-745-237A-4248 48 5 US-09-745-237A-4248 48 7 US-10-156-306-7370 40 7 US-10-109-349A-102 48 5 US-09-745-237A-4318 50 1 PCT-US02-12405-476 50 7 US-10-104-094-33 50 7 US-10-144-094-3 5 US-09-747-774A-80 45 5 US-09-747-774A-80 47 7 US-10-170-097-971 48 5 US-09-745-237A-4015 48 7 US-10-156-306-7410 48 7 US-10-156-306-7410 48 7 US-10-156-306-7410 48 7 US-10-156-306-7410 48 7 US-10-156-306-7653 5 US-09-718-331A-1013
tment of Alzheimer's Disease	Length 41; Indels 0; Gaps 0; ;73	SHISMS		Sequence 5, Appli Sequence 402, App Sequence 402, App Sequence 4248, Ap Sequence 7370, Ap Sequence 102, App Sequence 476, App Sequence 476, App Sequence 80, Appl Sequence 80, Appl Sequence 246, App Sequence 971, App Sequence 971, App Sequence 4015, Ap Sequence 4113, Ap Sequence 4143, Ap Sequence 7653, Ap Sequence 7653, Ap Sequence 7653, Ap

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Sequence 268, Application PC/TUS0212063

Sequence 268, Application:
GENERAL INFORMATION:
APPLICANT: SCHERING CORPORATION
APPLICANT: SCHERING CORPORATION
APPLICANT: GENOME THERAPEUTICS CORPORATION
TITLE OF INVENTION: NOVEL HUMAN GENE RELATING TO RESPIRATORY DISEASES,
TITLE OF INVENTION: OBESITY, AND INFLAMMATORY BOWEL DISEASE
FILE REFERENCE: 2976-4039PC2
CURRENT APPLICATION NUMBER: PCT/US02/12063
CURRENT APPLICATION NUMBER: PCT/US02/12063
CURRENT FILING DATE: 2002-04-15
PRIOR APPLICATION ONUMBER: 09/834,597
PRIOR FILING DATE: 2001-04-13
NUMBER OF SEQ ID NOS: 420
SOFTWARE: Patentin Ver. 2.1
SOFTWARE: Patentin Ver. 2.1
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PCT-US02-12063-268
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/980,217
CURRENT FILING DATE: 2002-05-06
PRIOR APPLICATION NUMBER: PCT/GB00/02072
PRIOR FILING DATE: 2000-05-30
PRIOR APPLICATION NUMBER: GB 9912563.5
PRIOR FILING DATE: 1999-05-28
NUMBER OF SEQ ID NOS: 52
SOFTWARE: Patentin Ver: 2.1
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US-09-980-217-51/c
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OTHER INFORMATION:
PCT-US02-12063-268
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APPLICANT: BLOTICA Technology Limited
APPLICANT: Leadlay, Peter F
APPLICANT: Staunton, James
APPLICANT: Staunton, James
APPLICANT: Oliynyk, Marko
TITLE OF INVENTION: Polyketides and their synthesis
FILE REFERENCE: IS/BP5858469
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 1.1%;
Best Local Similarity 63.6%;
Matches 28; Conservative
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Best Local Similarity 76.5%;
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Query Match
Best Local Similarity
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LENGTH: 50
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                  TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
                                                                                                                                              FEATURE:
                                                                                                                                                                                                     ENGTH: 41
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                                                                                                 Description of Artificial Sequence: Sequence containing SNP
1.1%;
82.8%;
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Pred. No. 4.1e+05;
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Pred. No. 3.
  Score
Pred.
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  N 21
4.4e+05;
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                     DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-10-126-022-268
                                                                                                                                                              ; OTHER INFORMATION: Description of Artificial Sequence: US-10-156-306-7580
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-10-156-306-7580
                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: RIBOZYME Pharmaccuticals, Inc.
APPLICANT: RIBOZYME Pharmaccuticals, Inc.
APPLICANT: RCSwiggen, James
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Rel
TITLE OF INVENTION: Levels of IKK-Gamma and PKR
FILE REPERINCE: MBHB01-664-A (400/050)
CURRENT APPLICATION NUMBER: US/10/156,306
CURRENT FILING DATE: 2002-05-28
NUMBER OF SEQ ID NOS: 8013
SOFTWARE: Patentin version 3.0
SEQ ID NO 7580
LENGTH: 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 09/548,797
PRIOR FILING DATE: 2000-04-13
NUMBER OF SEQ ID NOS: 420
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 268
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 82.8
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 268, Application US/10126022 GENERAL INFORMATION:
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                                                                     Query Match
Best Local Similarity
Matches 29; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: NOVEL HUMAN GENE RELATING TO RESPIRATORY DISEASES, TITLE OF INVENTION: OBESITY, AND INFLAMMATORY BOWEL DISEASE FILE REFERENCE: 2976-4039US2 CURRENT APPLICATION NUMBER: US/10/126,022 CURRENT FILING DATE: 2002-04-19 PRIOR APPLICATION NUMBER: 09/834,597 PRIOR RILING DATE: 2001-04-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: KEITH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 41
                                                                                                                                                                                                                 ORGANISM: Artificial Sequence
                                                                                                                                                                                                FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          624 acagcagctgggcagccaaggcccctggg
             137 ggctgggggtaggggactccctacaggcacgcagccctgagaccgcca 184
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                                                                   Score 20.8; D. Pred. No. 4.8e 2; Mismatches
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Pred. No.
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                                                                       .8e+05;
es 17;
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                                                                                                       Length 48;
                                                                       Indels
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US-09-978-917A-25/c; Sequence 25, Application US/09978917A; GENERAL INFORMATION:
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; OTHER INFORMATION: Accession number cg43955639
US-09-718-321A-1048
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RESULT
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CURRENT FILING DATE: 2002-06-20
PRIOR APPLICATION NUMBER: 60/163,783
PRIOR FILING DATE: 1999-11-24
PRIOR SEQ ID NOS: 1468
SOFTWARE: CuraGen Patent Formatter Version 0.9
SEQ ID NO 1048
LENGTH: 50
                                                                                                                                                Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 48
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 25
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Maxygen Aps; Maxygen Holdings
TITLE OF INVENTION: Protein C or activated protein C-like molecules
FILE REFERENCE: 0219us310 - protein C
CURRENT APPLICATION NUMBER: US/09/978,917A
CURRENT FILING DATE: 2001-10-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Shimkets, Richard A.
APPLICANT: Leach, Martin D.
TITLE OF INVENTION: USE THEREOF
TITLE OF INVENTION: USE THEREOF
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                                                                                                                                                                                                                                                                         TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: allele
LOCATION: (26)...(0)
OTHER INFORMATION: single nucleotide polymorphism
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OTHER INFORMATION: Nucleotide deleted between bases 25 and 26
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                                                                            713 gccccagcgggcaccctggacttggtgcccggcgggctgacct 755
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                                                         46 GCCCGCAGGCCAGCTTGTTGTTGTCCAGCAGGACCACCT
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                                                                                                                                     Conservative
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Pred. No. 4.8e+05;
0; Mismatches 17;
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Pred. No. 5.2e+05;
                                                                                                                                   Mismatches
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US-60-384-980-69/c
Sequence 69, Application US/60384980
GENERAL INFORMATION:
APPLICANT: Bristol-Myers Squibb Company
TITLE OF INVENTION: HUMAN SINGLE NUCLEOTIDE POLYMORPHISMS
FILE REFERENCE: D0211 PSP
CURRENT APPLICATION NUMBER: US/60/384,980
CURRENT FILING DATE: 2002-06-03
NUMBER OF SEQ ID NOS: 1219
SOFTMARE: Patentin version 3.1
SEQ ID NO 69
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SEQ ID NO 25
LENGTH: 49
TYPE: DNA
ORGANISM: Artificial Sequence
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GENERAL INFORMATION:
APPLICANT: MAXYGEN Aps; MAXYGEN Holdings
FITLE OF INVENTION: Protein C or activated
FILE REFERENCE: 0219us410 - protein C
FULE REFERENCE: 0219us410 - protein C
CURRENT APPLICATION UNMERS: US/09/997,623
CURRENT FILING DATE: 2001-11-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; OTHER INFORMATION: Description of Artificial Sequence: US-09-745-237A-3833
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                                                                                                                                                                                                                                                             RESULT 11
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APPLICANT: McSwiggen, Jim
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Method and Reagent for the Treatment of Alzheimer's Disease
FILE REFERENCE: 400/007 (MBHB00-918-A)
CURRENT FILING DATE: 2002-04-15
NUMBER OF SEQ ID NOS: 4550
SOFTWARE: Patentin version 3.0
SEQ ID NO 3833
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APPLICANT: Ribozyme Pharmaceuticals, Inc.
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Best Local Similarity
Matches 28; Conserv
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Best Local Similarity
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PRIOR FILING DATE: 2001-10-17
NUMBER OF SEQ ID NOS: 48
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: RNA
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Pred. No. 5.7e+05;
2; Mismatches 16
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Pred. No. 5.2e+05;
0; Mismatches 14;
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APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim
APPLICANT: McSwiggen, Jim
FILE OF INVENTION: Method and Reagent for the Treatment of
FILE REFERENCE: 400/007 (MBHB00-918-A)
CURRENT APPLICATION NUMBER: US/09/745,237A
CURRENT FILING DATE: 2002-04-15
NUMBER OF SEQ ID NOS: 4550
SOFTWARE: PatentIn version 3.0
SEQ ID NO 3872
LENGTH: 48
TYPE: RNA
ORGANISM: Artificial Sequence
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-60-384-980-69
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1257
LENGTH: 50
TYPE: DNA
CRGANIZM: Homo sapiens
US-10-099-926-1257
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                                                                                             ; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence:
US-09-745-237A-3872
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US-10-099-926-1257/c
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APPLICANT: Meaghe
APPLICANT: Xu, Ji
APPLICANT: Secris
APPLICANT: Jiang
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Best Local Similarity 68.3%;
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Best Local Similarity
                                     Query Match
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APPLICANT: Jiang, Yuqiu
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF COLON CANCER
FILE REFERENCE: 210121.547C2
CURRENT APPLICATION NUMBER: US/10/099,926
CURRENT FILING DATE: 2002-03-17
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   Local Similarity
les 27; Conserv
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Xu, Jiangchun
   Conservative
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                   1.18;
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Score 20; DB 5; Le
Pred. No. 6.8e+05;
Prematches 15;
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Pred. No. 6.2e+05;
0; Mismatches 18;
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Pred. No. 6.2e+05;
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                                     Length 48
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                                                                                                                 Enzymatic Nucleic Acid
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PRIOR APPLICATION NUMBER: PCT/EPO 0/12419
PRIOR FILING DATE: 2000-12-08
PRIOR APPLICATION NUMBER: US 60/169,653
PRIOR FILING DATE: 1999-12-08
NUMBER OF SEQ ID NOS: 85
SOFTWARE: FASTSEQ FOR Windows Version 4.0
SEQ ID NO 45
LENGTH: 44
TYPE: DNA
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US-09-745-237A-3877
; Sequence 3877, Application US/09745237A
; GENERAL INFORMATION:
; GENERAL INFORMATION:
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; OTHER INFORMATION: Description of Artificial Sequence:
US-09-745-237A-3877
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LENGTH: 48

TYPE: RNA

ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
Matches 27; Conserv
                                                                              Query Match
Best Local Similarity
Matches 27; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Method and Reagent for the Treatment of Alzheimer's Disease
FILE REFERENCE: 400/007 (MBHB00-918-A)
CURRENT APPLICATION NUMBER: US/09/745,237A
CURRENT FILING DATE: 2002-04-15
NUMBER OF SEQ ID NOS: 4550
SOFTMARE: Patentin version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Steinberger, Peter
TITLE OF INVENTION: METHODS AND COMPOSITIONS USEFUL FOR
TITLE OF INVENTION: INHIBITING CCR5-DEPENDENT INFECTION OF CELLS BY HIV-1
FILE REFERENCE: TSRI 728.1
CURRENT APPLICATION NUMBER: US/09/913,238
CURRENT FILING DATE: 2002-03-05
CURRENT FILING DATE: 2002-03-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Barbas, Carlos F. APPLICANT: Steinberger, Pet
                                                                                                                                                                                                                            ORGANISM: Unknown FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                139 ctgggggtaggggactccctacaggcacgcagccctgagaccgc 182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        138 gctgggggtaggggactccctacaggcacgcacgccctgagaccg 181
                     358 cagctccctgggcaccaggcagctaacagacacagccgc 396
39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3 ccggggggaggaaacucccuucaaggacaucguccgggagccgc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2 gcagggggggggagaacucccuucaaggacaucguccgggagccg 45
CGGCCGCCTGGGCCCCTCGCAGCTAGCAAGCTCCTCCGC 1
                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1.1%;
                                                                                                    1.0%;
                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 20; DB 5;
Pred. No. 6.8e+05;
2; Mismatches 15
                                                                              Score 19.8; DB 5;
Pred. No. 7.4e+05;
0; Mismatches 12
                                                                                                                        DB 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15;
                                                                                  12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 48;
                                                                                                                        Length 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 46
                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Enzymatic Nucleic Acid
                                                                                  0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
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Search completed: August

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Job time: 7651 sec

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